



IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

APPLICANT: Prayaga et al.

SERIAL NUMBER: 09/679,740

EXAMINER: Not Yet Assigned

FILING DATE: October 5, 2000

ART UNIT: 1645

FOR: Endozepine-Like Polypeptides and Polynucleotides Encoding Same

Mail Stop Missing Parts
Commissioner for Patents
P.O. Box 1450
Alexandria, VA 22313-1450

**RESPONSE TO NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT
APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID
SEQUENCE DISCLOSURES**

In response to the notice to comply with requirements for patent applications containing nucleotide sequence and/or amino acid sequence disclosures, mailed February 14, 2002, in the above-identified application, Applicants submit a substitute paper copy, and a computer readable form of the Sequence Listing. Also enclosed are a Statement in Support of Computer Readable Form Submission, a Supplemental Preliminary Amendment and a copy of the Notice to Comply. Applicants also enclose a copy of a Petition Under 37 C.F.R. § 1.181 to Withdraw Holding of Abandonment that is being filed simultaneously with this Response.

Please charge any fees that may be due, or credit any overpayment to Deposit Account No. 50-0311, Reference No. 15966-575B.

Respectfully submitted,

Date of Deposit: July 22, 2004

Christina K. Rock, Reg. No. 45,899
Ivor R. Elrifi, Reg. No. 39,529
Attorney for Applicants
c/o MINTZ, LEVIN
Tel: (617) 542-6000
Fax: (617) 542-2241
Customer No.: 30623



SEQUENCE LISTING

<110> Prayaga, Sudhirdas K
Shimkets, Richard A
Majumder, Kumud
Eisen, Andrew
Vernet, Corine
Spaderna, Steven K

<120> ENDOZEPINE-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES
ENCODING SAME

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<141> 2000-10-05

<150> 60/157,786
<151> 1999-10-05

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<151> 1999-11-09

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<170> PatentIn Ver. 2.1

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 cccgtgagcg atcaggagaa gctgctgggc tacggcttgt acaaacaggc caccaggggc 180
 gactgcgaca tccccggccc tccggcctca gacgtgagag ccagggccaa gtgggagggt 240
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 35 40 45
 Leu Val Tyr Gly Leu Tyr Lys Gln Ala Thr Gln Gly Asp Cys Asp Ile
 50 55 60
 Pro Gly Pro Pro Ala Ser Asp Val Arg Ala Arg Ala Lys Trp Glu Ala
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 Ala Ala Lys Val Glu Glu Leu Thr Lys Lys Glu
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 gatgatgaag aactgaaaga actttatggg ctttacaaac aagctgtaat tggaaacatt 180
 aatattgagt gttcagaaat gctagaatta aaaggcaagg ccaaattggga agcacagaac 240
 ccccaaaaag gattgtcaga ggaagatatg atgcgtgcct ttattttctaa agccgaagag 300

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 35 40 45
 Leu Glu Leu Lys Gly Lys Ala Lys Trp Glu Ala Gln Asn Pro Gln Lys
 50 55 60
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 caagccagca gatgatgaga tgcggttcct ttacggccac taaaacgag cgactgtagg 300
 caacataaag acagaacggc cagggatggt ggacttcaag ggcaaagcca agtgggatcc 360
 ctggaattta gtgaaagggg ctgccaggga agatcccatg aaagctaaag cttacgtcaa 420
 aaaagtagaa gagttaaaga aaaaattcag aatacgagag actggaattg ttgccagcca 480
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His Leu Ser Trp Glu Glu Lys Lys Lys Lys Lys Arg Cys Ala Gly Ile
 35 40 45
 Lys His Phe Lys Thr Lys Pro Ala Asp Asp Glu Met Arg Phe Leu Tyr
 50 55 60
 Gly His Tyr Lys Arg Ala Thr Val Gly Asn Ile Lys Thr Glu Arg Pro
 65 70 75 80
 Gly Met Val Asp Phe Lys Gly Lys Ala Lys Trp Asp Pro Trp Asn Leu
 85 90 95
 Val Lys Gly Ala Ala Arg Glu Asp Pro Met Lys Ala Lys Ala Tyr Val
 100 105 110
 Lys Lys Val Glu Glu Leu Lys Lys Lys Phe Arg Ile Arg Glu Thr Gly
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 Ile Val Ala Ser His Ala Phe Val Leu Asn
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 ctatgggctt tacaacaag caatagttgg agacattaat attgcgtgtc caggaatgct 180
 agatttaaaa ggcaaagcca aatgggaagc atggaacctc aaaaaagggg tgtcgacgga 240
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 gaatacagca 310

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 Glu Leu Lys Glu Leu Tyr Gly Leu Tyr Lys Gln Ala Ile Val Gly Asp
 35 40 45
 Ile Asn Ile Ala Cys Pro Gly Met Leu Asp Leu Lys Gly Lys Ala Lys
 50 55 60
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<212> DNA
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20 25 30
Leu Tyr Lys Gln Ala Ile Ile Gly Asp Ile Asn Ile Glu Tyr Leu Gly
35 40 45
Met Leu Asp Phe Lys Gly Lys Ala Lys Cys Ala Ala Trp Thr Leu Gln
50 55 60
Lys Arg Leu Ser Lys Glu Asp Ala Thr Ser Val Ser Ile Ser Lys Ala
65 70 75 80
Lys Glu Pro Ile Glu Lys
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 Tyr Gly Leu Tyr Lys Gln Ala Thr Gln Gly Asp Cys Asp Ile Pro Gly
 35 40 45
 Pro Pro Ala Ser Asp Val Arg Ala Arg Ala Lys Trp Glu Ala Trp Ser
 50 55 60
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Phe Lys

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<213> Homo sapiens

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Leu Lys Gly Lys
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<210> 20

<211> 18

<212> PRT

<213> Homo sapiens

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Leu Lys

<210> 21

<211> 20

<212> PRT

<213> Homo sapiens

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      35                      40                      45

Val Lys Val Ile Gln Ser Leu Pro Lys Asn Gly Ser Phe Gln Pro Thr
      50                      55                      60

Asn Glu Met Met Leu Lys Phe Tyr Ser Phe Tyr Lys Gln Ala Thr Glu
      65                      70                      75                      80

Gly Pro Cys Lys Leu Ser Arg Pro Gly Phe Trp Asp Pro Ile Gly Arg
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Tyr Lys Trp Asp Ala Trp Ser Ser Leu Gly Asp Met Thr Lys Glu Glu
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Ala Met Ile Ala Tyr Val Glu Glu Met Lys Lys Ile Ile Glu Thr Met
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Pro Met Thr Glu Lys Val Glu Glu Leu Leu Arg Val Ile Gly Pro Phe
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Tyr Glu Ile Val Glu Asp Lys Lys Ser Gly Arg Ser Ser Asp Ile Thr
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Ser Ser Asp Ser Gly Ala Glu Ser Glu Glu Glu Glu Ala Gln Glu Glu
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Ser Ala Asp His Lys Asn Leu Glu Val Ile Val Thr Asn Gly Tyr Asp
225 230 235 240
Lys Asp Gly Phe Val Gln Asp Ile Gln Asn Asp Ile His Ala Ser Ser
245 250 255
Ser Leu Asn Gly Arg Ser Thr Glu Glu Val Lys Pro Ile Asp Glu Asn
260 265 270
Leu Gly Gln Thr Gly Lys Ser Ala Val Cys Ile His Gln Gly Ile Asn
275 280 285
Asp Asp His Val Glu Asp Val Thr Gly Ile Gln His Leu Thr Ser Asp
290 295 300
Ser Asp Ser Glu Val Tyr Cys Asp Ser Met Glu Gln Phe Gly Gln Glu
305 310 315 320
Glu Ser Leu Asp Ser Phe Thr Ser Asn Asn Gly Pro Phe Gln Tyr Tyr
325 330 335
Leu Gly Gly His Ser Ser Gln Pro Met Glu Asn Ser Gly Phe Arg Glu
340 345 350
Asp Ile Gln Val Pro Pro Gly Asn Gly Asn Ile Gly Asn Met Gln Val
355 360 365
Val Ala Val Glu Gly Lys Gly Glu Val Lys His Gly Gly Glu Asp Gly
370 375 380
Arg Asn Asn Ser Gly Ala Pro His Arg Glu Lys Arg Gly Gly Glu Thr
385 390 395 400
Asp Glu Phe Ser Asn Val Arg Arg Gly Arg Gly His Arg Met Gln His
405 410 415
Leu Ser Glu Gly Thr Lys Gly Arg Gln Val Gly Ser Gly Gly Asp Gly
420 425 430
Glu Arg Trp Gly Ser Asp Arg Gly Ser Arg Gly Ser Leu Asn Glu Gln
435 440 445
Ile Ala Leu Val Leu Met Arg Leu Gln Glu Asp Met Gln Asn Val Leu
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Gln Arg Leu Gln Lys Leu Glu Thr Leu Thr Ala Ala Lys Ser Ser Thr
465 470 475 480
Ser Thr Leu Gln Thr Ala Pro Gln Pro Thr Ser Ser Gln Arg Pro Ser
485 490 495
Trp Trp Pro Phe Glu Met Ser Pro Gly Val Leu Thr Phe Ala Ile Ile
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Arg Arg
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Gln Ala Thr Val His Asp Leu Asn Thr Glu Trp Pro Arg Met Leu Asp
35 40 45

Leu Lys Gly Lys Ala Lys Gln Asp Ala Trp Asn Glu Leu Lys Asp Thr
50 55 60

Ala Lys Glu Asp Ala Val Lys Ala Asp Ile Asn Lys Val Glu Glu Arg
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Asn Lys Lys Tyr Arg Ile
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Leu Lys Gly Lys
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 aaatacggga tatga 315

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 20 25 30

Leu Lys Thr Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His
 35 40 45

Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met
 50 55 60

Leu Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys
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Glu Leu Lys Lys Lys Tyr Gly Ile
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Phe Thr Gly Lys
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<210> 31
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<210> 32
 <211> 359
 <212> PRT
 <213> Homo sapiens

<400> 32
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 20 25 30

Leu Tyr Lys Gln Ala Thr Glu Gly Pro Cys Asn Met Pro Lys Pro Gly
 35 40 45

Val Phe Asp Leu Ile Asn Lys Ala Lys Trp Asp Ala Trp Asn Ala Leu
 50 55 60

Gly Ser Leu Pro Lys Glu Ala Ala Arg Gln Asn Tyr Val Asp Leu Val

65		70		75		80
Ser Ser Leu Ser Pro Ser Leu Glu Ser Ser Ser Gln Val Glu Pro Gly						
		85		90		95
Thr Asp Arg Lys Ser Thr Gly Phe Glu Thr Leu Val Val Thr Ser Glu						
	100		105		110	
Asp Gly Ile Thr Lys Ile Met Phe Asn Arg Pro Lys Lys Lys Asn Ala						
	115		120		125	
Ile Asn Thr Glu Met Tyr His Glu Ile Met Arg Ala Leu Lys Ala Ala						
	130		135		140	
Ser Lys Asp Asp Ser Ile Ile Thr Val Leu Thr Gly Asn Gly Asp Tyr						
145		150		155		160
Tyr Ser Ser Gly Asn Asp Leu Thr Asn Phe Thr Asp Ile Pro Pro Gly						
	165		170			175
Gly Val Glu Glu Lys Ala Lys Asn Asn Ala Val Leu Leu Arg Glu Phe						
	180		185		190	
Val Gly Cys Phe Ile Asp Phe Pro Lys Pro Leu Ile Ala Val Val Asn						
	195		200		205	
Gly Pro Ala Val Gly Ile Ser Val Thr Leu Leu Gly Leu Phe Asp Ala						
	210		215		220	
Val Tyr Ala Ser Asp Arg Ala Thr Phe His Thr Pro Phe Ser His Leu						
225		230		235		240
Gly Gln Ser Pro Glu Gly Cys Ser Ser Tyr Thr Phe Pro Lys Ile Met						
	245		250			255
Ser Pro Ala Lys Ala Thr Glu Met Leu Ile Phe Gly Lys Lys Leu Thr						
	260		265		270	
Ala Gly Glu Ala Cys Ala Gln Gly Leu Val Thr Glu Val Phe Pro Asp						
	275		280		285	
Ser Thr Phe Gln Lys Glu Val Trp Thr Arg Leu Lys Ala Phe Ala Lys						
	290		295		300	
Leu Pro Pro Asn Ala Leu Arg Ile Ser Lys Glu Val Ile Arg Lys Arg						
305		310		315		320
Glu Arg Glu Lys Leu His Ala Val Asn Ala Glu Glu Cys Asn Val Leu						
	325		330		335	
Gln Gly Arg Trp Leu Ser Asp Glu Cys Thr Asn Ala Val Val Asn Phe						
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<210> 33
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 <213> Homo sapiens

<400> 33
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 Leu Ile Asn Lys
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<210> 34
 <211> 1574
 <212> DNA
 <213> Homo sapiens

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 ganaggacgt ccagcgtacg tngccccg cttccccgcc ggcgcagagc aggcctcaca 180
 gaatcgcacg ccgctggcac gcacgccgcc ccgccccac ggcccagcgc cagcgcgccc 240
 cgcgtcgcac gcatcccggc ctcaactgcc ctgcactcct gttccgttgg aggggcctga 300
 ggcgagcctg agcgcgctgt tggccggagg aagccggaga gaccgggtcg actgggcaga 360
 gcggcagagg gtcgaggagc ctgctctgca cgcccaggga gtagaagtgg gcaggagaca 420
 gggtcacgtg agggagcgcg ccgcgactga gcttgggtcc gactggagct caggctcgcg 480
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 aagagggtag ggctctactt cactgggcct gtgatcgagg acataaggaa ctagtcacag 1200
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<210> 35
 <211> 282
 <212> PRT
 <213> Homo sapiens

<400> 35

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			20					25					30			
Ser	Pro	Glu	Ile	Glu	Glu	Thr	Ser	Cys	Leu	Ala	Glu	Leu	Phe	Glu	Lys	
		35					40					45				
Ala	Ala	Ala	His	Leu	Gln	Gly	Leu	Ile	Gln	Val	Ala	Ser	Arg	Glu	Gln	
	50					55					60					
Leu	Leu	Tyr	Leu	Tyr	Ala	Arg	Tyr	Lys	Gln	Val	Lys	Val	Gly	Asn	Cys	
65					70					75					80	
Asn	Thr	Pro	Lys	Pro	Ser	Phe	Phe	Asp	Phe	Glu	Gly	Lys	Gln	Lys	Trp	
				85					90					95		
Glu	Ala	Trp	Lys	Ala	Leu	Gly	Asp	Ser	Ser	Pro	Ser	Gln	Ala	Met	Gln	
			100					105					110			
Glu	Tyr	Ile	Ala	Val	Val	Lys	Lys	Leu	Asp	Pro	Gly	Trp	Asn	Pro	Gln	
	115						120					125				
Ile	Pro	Glu	Lys	Lys	Gly	Lys	Glu	Ala	Asn	Thr	Gly	Phe	Gly	Gly	Pro	
	130					135					140					
Val	Ile	Ser	Ser	Leu	Tyr	His	Glu	Glu	Thr	Ile	Arg	Glu	Glu	Asp	Lys	
145					150					155					160	
Asn	Ile	Phe	Asp	Tyr	Cys	Arg	Glu	Asn	Asn	Ile	Asp	His	Ile	Thr	Lys	
				165					170					175		
Ala	Ile	Lys	Ser	Lys	Asn	Val	Asp	Val	Asn	Val	Lys	Asp	Glu	Glu	Gly	
			180					185					190			
Arg	Ala	Leu	Leu	His	Trp	Ala	Cys	Asp	Arg	Gly	His	Lys	Glu	Leu	Val	
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Thr	Val	Leu	Leu	Gln	His	Arg	Ala	Asp	Ile	Asn	Cys	Gln	Asp	Asn	Glu	
	210					215					220					
Gly	Gln	Thr	Ala	Leu	His	Tyr	Ala	Ser	Ala	Cys	Glu	Phe	Leu	Asp	Ile	
225					230					235					240	
Val	Glu	Leu	Leu	Leu	Gln	Ser	Gly	Ala	Asp	Pro	Thr	Leu	Arg	Asp	Gln	
				245					250					255		
Asp	Gly	Cys	Leu	Pro	Glu	Glu	Val	Thr	Gly	Cys	Lys	Thr	Val	Ser	Leu	
			260					265					270			
Val	Leu	Gln	Arg	His	Thr	Thr	Gly	Lys	Ala							
		275					280									

<210> 36
<211> 20
<212> PRT
<213> Homo sapiens

<400> 36
Gln Val Lys Val Gly Asn Cys Asn Thr Pro Lys Pro Ser Phe Phe Asp
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Phe Glu Gly Lys
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<210> 37
<211> 20
<212> PRT
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<223> wherein Xaa is Leu or Phe

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 <222> (20)
 <223> wherein Xaa is Lys or Arg

 <400> 37
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 1 5 10 15

 Xaa Lys Gly Xaa
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<210> 38
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<221> VARIANT

<222> (18)

<223> wherein Xaa is Lys or Ile

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<222> (20)

<223> wherein Xaa is Lys or Arg

<400> 38

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1

5

10

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Xaa Xaa Gly Xaa

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<210> 39

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 or Met

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 or Met

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 Xaa Ile Xaa Xaa
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<210> 40
 <211> 20
 <212> PRT
 <213> Homo sapiens

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 <223> wherein Xaa is Thr, Lys or Glu

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 1 5 10 15

Phe Xaa Gly Lys
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<210> 41
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 <212> PRT
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1 5 10 15

Xaa Xaa Gly Lys
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<210> 42

<211> 20

<212> PRT
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Xaa Xaa Gly Xaa
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<223> wherein Xaa is Gly or Ala

<220>
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<223> wherein Xaa is Lys or Arg

<400> 43

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1 5 10 15

Xaa Xaa Xaa Xaa
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<210> 44

<211> 20

<212> PRT

<213> Homo sapiens

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 1 5 10 15

 Xaa Xaa Gly Lys
 20

 <210> 45
 <211> 20
 <212> PRT
 <213> Homo sapiens

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<220>
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 1 5 10 15

 Phe Xaa Gly Lys
 20

<210> 46
 <211> 687
 <212> DNA
 <213> Homo sapiens

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 ccatgtgcc aagtggagttc gagctgcgcg gccctcaagc agctgaaggg tcccgtgagc 240
 gatcaggaga agctgctggt ctacggcttg taaaaacagg ccaccaggcg cgactgcgac 300
 atccccggcc ctccggcctc agacgtgaga gccagggcca agtgggaggc ttggagcgcg 360
 aacaaagggg cgtccaagat ggacgccatg aggggctacg cggccaaagt ggaggagctg 420
 acgaagaagg aagtggggggg cgtggagcgc gaacaaaggg gcgtgcaaga tggacgccat 480
 gaggggctac gcgcccaaag tggaggagct gacgaagaag gaaggcgctc caagatggac 540
 gccatgaggg gctacgcggc caaagtggag gagctgacga agaaggaagt ggggggcgtg 600
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 gagatgagga agaaggaggc tggctga 687

<210> 47
 <211> 228
 <212> PRT
 <213> Homo sapiens

<400> 47

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Arg Thr Ser Ala Pro Ser Ala Gln Arg Arg Leu Pro Ala Glu Pro Ser
          35              40              45

His Gln Pro Ser Ala Pro Gly Thr Ala Ser Thr Thr Pro Cys Ala Lys
 50              55              60

Trp Ser Ser Ser Cys Ala Ala Leu Lys Gln Leu Lys Gly Pro Val Ser
 65              70              75              80

Asp Gln Glu Lys Leu Leu Val Tyr Gly Leu Tyr Lys Gln Ala Thr Gln
          85              90              95

Gly Asp Cys Asp Ile Pro Gly Pro Pro Ala Ser Asp Val Arg Ala Arg
100              105              110

Ala Lys Trp Glu Ala Trp Ser Ala Asn Lys Gly Ala Ser Lys Met Asp
115              120              125

Ala Met Arg Gly Tyr Ala Ala Lys Val Glu Glu Leu Thr Lys Lys Glu
130              135              140

Val Gly Gly Val Glu Arg Glu Gln Arg Gly Val Gln Asp Gly Arg His
145              150              155              160

Glu Gly Leu Arg Gly Gln Ser Gly Gly Ala Asp Glu Glu Gly Arg Ala
          165              170              175

Ser Lys Met Asp Ala Met Arg Gly Tyr Ala Ala Lys Val Glu Glu Leu
180              185              190

Thr Lys Lys Glu Val Gly Gly Val Glu Arg Glu Gln Arg Gly Val Gln
195              200              205

Asp Gly Arg His Glu Gly Leu Arg Gly Gln Ser Glu Glu Met Arg Lys
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Lys Glu Ala Gly
225

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 <212> DNA
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gccttccggc agagccctcc caccagccct cagcttctag caccagggac cgctccacc 180
accccatgtg ccaagtggag ttcgagctgc gcggccctca agcagctgaa ggggtcccgtg 240
agcgatcagg agaagctgct ggtctacggc ttgtacaaac aggccaccca gggcgactgc 300
gacatccccg gccctccggc ctcagacgtg agagccaggg ccaagtggga ggcttggagc 360
gcgaaaaaag gggcggtccaa gatggacgcc atgaggggct acgcgggcaa agtggaggag 420
ctgacgaaga aggaagtggg gggcggtggag cgcgaaacaa gggggcgtgca agatggacgc 480
catgaggggc tacgcggcca aagtggagga gctgacgaag aaggaagtgg ggggcgtgga 540
gcgcgaacaa agggggcgtcc aagatggacg ccatga 576

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<210> 49
<211> 191
<212> PRT
<213> Homo sapiens

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<400> 49
Met Gly Asp Ala Gly Ala Thr Ala Ala Ala Leu Arg Pro Ala His Asn
  1             5             10             15

Leu Arg Pro Ala Pro Pro Thr Ala Ser Ala Ala His Ala Ser Pro His
      20             25             30

Glu Arg Ala Arg Gln Ala Ser Arg Ala Phe Arg Gln Ser Pro Pro Thr
  35             40             45

Ser Pro Gln Leu Leu Ala Pro Gly Thr Ala Ser Thr Thr Pro Cys Ala
  50             55             60

Lys Trp Ser Ser Ser Cys Ala Ala Leu Lys Gln Leu Lys Gly Pro Val
  65             70             75             80

Ser Asp Gln Glu Lys Leu Leu Val Tyr Gly Leu Tyr Lys Gln Ala Thr
      85             90             95

Gln Gly Asp Cys Asp Ile Pro Gly Pro Pro Ala Ser Asp Val Arg Ala
  100            105            110

Arg Ala Lys Trp Glu Ala Trp Ser Ala Lys Lys Gly Ala Ser Lys Met
  115            120            125

Asp Ala Met Arg Gly Tyr Ala Ala Lys Val Glu Glu Leu Thr Lys Lys
  130            135            140

Glu Val Gly Gly Val Glu Arg Glu Gln Arg Gly Val Gln Asp Gly Arg
  145            150            155            160

His Glu Gly Leu Arg Gly Gln Ser Gly Gly Ala Asp Glu Glu Gly Ser
      165            170            175

Gly Gly Arg Gly Ala Arg Thr Lys Gly Arg Pro Arg Trp Thr Pro
  180            185            190

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<210> 50

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<211> 294
 <212> DNA
 <213> Homo sapiens

<400> 50
 gctgcgccca ccatgtccct gcaggctgat tttgacatgg tcacagaaga tgtgaggaag 60
 ctgaaaacaa gaccagatga tgaagaactg aaagaacttt atgggcttta caaacaagct 120
 gtaattggaa acattaatat tgagtgttca gaaatgctag aattaaaagg caaggccaaa 180
 tgggaagcac agaaccacca aaaaggattg tcagaggaag atatgatgcg tgcctttatt 240
 tctaaagccg aagagctgat agaaaaatat ggaatttaga ataaagcata tgat 294

<210> 51
 <211> 293
 <212> DNA
 <213> Homo sapiens

<400> 51
 gctgaatcaa ccatgtcacc ccaggcagat tttgacaaag cagcagggga tgtaaagaaa 60
 ttgaaaacaa aaccaactga cgatgaactg aaggaactgt acggactcta caagcagtcc 120
 actgttgggg acataaatat agagtgtcct ggcattgctag atctgaaggg caaggccaag 180
 tgggacgcat ggaacctaaa gaaaggcttg tctaaggaag atgcgatgag cgcttatgtt 240
 tctaaagccc atgagctgat agaaaaatat ggcctgtaac aaggtcgcg gat 293

<210> 52
 <211> 85
 <212> PRT
 <213> Homo sapiens

<400> 52
 Gln Ala Asp Phe Asp Met Val Thr Glu Asp Val Arg Lys Leu Lys Thr
 1 5 10 15
 Arg Pro Asp Asp Glu Glu Leu Lys Glu Leu Tyr Gly Leu Tyr Lys Gln
 20 25 30
 Ala Val Ile Gly Asn Ile Asn Ile Glu Cys Ser Glu Met Leu Glu Leu
 35 40 45
 Lys Gly Lys Ala Lys Trp Glu Ala Gln Asn Pro Gln Lys Gly Leu Ser
 50 55 60
 Glu Glu Asp Met Met Arg Ala Phe Ile Ser Lys Ala Glu Glu Leu Ile
 65 70 75 80
 Glu Lys Tyr Gly Ile
 85

<210> 53
 <211> 85
 <212> PRT
 <213> Homo sapiens

<400> 53
 Gln Ala Asp Phe Asp Glu Ala Ala Glu Glu Val Lys Lys Leu Lys Thr

1	5	10	15
Arg	Pro	Thr	Asp
	Glu	Glu	Leu
	Lys	Glu	Leu
	Tyr	Gly	Phe
	Tyr	Lys	Gln
	20	25	30
Ala	Thr	Val	Gly
	Asp	Ile	Asn
	Ile	Glu	Cys
	Pro	Gly	Met
	Leu	Asp	Leu
	35	40	45
Lys	Gly	Lys	Ala
	Lys	Trp	Glu
	Ala	Trp	Asn
	Leu	Lys	Lys
	Gly	Ile	Ser
	50	55	60
Lys	Glu	Asp	Ala
	Met	Asn	Ala
	Tyr	Ile	Ser
	Lys	Ala	Lys
	Thr	Met	Val
	65	70	75
Glu	Lys	Tyr	Gly
	Ile		
	85		

<210> 54
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 54
Ser
Gln
Ala
Glu
Phe
Glu
Lys
Ala
Ala
Glu
Glu
Val
Lys
Asn
Leu
Lys
1
5
10
15
Thr
Lys
Pro
Ala
Asp
Asp
Glu
Met
Leu
Phe
Ile
Tyr
Ser
His
Tyr
Lys
20
25
30
Gln
Ala
Thr
Val
Gly
Asp
Ile
Asn
Thr
Glu
Arg
Pro
Gly
Ile
Leu
Asp
35
40
45
Leu
Lys
Gly
Lys
Ala
Lys
Trp
Asp
Ala
Trp
Asn
Gly
Leu
Lys
Gly
Thr
50
55
60
Ser
Lys
Glu
Asp
Ala
Met
Lys
Ala
Tyr
Ile
Asn
Lys
Val
Glu
Glu
Leu
65
70
75
80
Lys
Lys
Lys
Tyr
Gly
Ile
85

<210> 55
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 55
Ser
Gln
Ala
Glu
Phe
Asp
Lys
Ala
Ala
Glu
Glu
Val
Lys
His
Leu
Lys
1
5
10
15
Thr
Lys
Pro
Ala
Asp
Glu
Glu
Met
Leu
Phe
Ile
Tyr
Ser
His
Tyr
Lys
20
25
30
Gln
Ala
Thr
Val
Gly
Asp
Ile
Asn
Thr
Glu
Arg
Pro
Gly
Met
Leu
Asp
35
40
45

Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys Gly Thr
50 55 60

Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asp Lys Val Glu Glu Leu
65 70 75 80

Lys Lys Lys Tyr Gly Ile
85

<210> 56
<211> 86
<212> PRT
<213> Homo sapiens

<400> 56
Ser Gln Ala Glu Phe Glu Lys Ala Ala Glu Glu Val Arg His Leu Lys
1 5 10 15

Thr Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His Tyr Lys
20 25 30

Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu Asp
35 40 45

Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys Gly Thr
50 55 60

Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asn Lys Val Glu Glu Leu
65 70 75 80

Lys Lys Lys Tyr Gly Ile
85

<210> 57
<211> 88
<212> PRT
<213> Homo sapiens

<400> 57
Met Ser Leu Gln Ala Asp Phe Asp Met Val Thr Glu Asp Val Arg Lys
1 5 10 15

Leu Lys Thr Arg Pro Asp Asp Glu Glu Leu Lys Glu Leu Tyr Gly Leu
20 25 30

Tyr Lys Gln Ala Val Ile Gly Asn Ile Asn Ile Glu Cys Ser Glu Met
35 40 45

Leu Glu Leu Lys Gly Lys Ala Lys Trp Glu Ala Gln Asn Pro Gln Lys
50 55 60

Gly Leu Ser Glu Glu Asp Met Met Arg Ala Phe Ile Ser Lys Ala Glu
65 70 75 80

Glu Leu Ile Glu Lys Tyr Gly Ile
85

<210> 58
<211> 82
<212> PRT
<213> Homo sapiens

<400> 58
Lys Arg Cys Ala Gly Ile Lys His Phe Lys Thr Lys Pro Ala Asp Asp
1 5 10 15
Glu Met Arg Phe Leu Tyr Gly His Tyr Lys Arg Ala Thr Val Gly Asn
20 25 30
Ile Lys Thr Glu Arg Pro Gly Met Val Asp Phe Lys Gly Lys Ala Lys
35 40 45
Trp Asp Pro Trp Asn Leu Val Lys Gly Ala Ala Arg Glu Asp Pro Met
50 55 60
Lys Ala Lys Ala Tyr Val Lys Lys Val Glu Glu Leu Lys Lys Lys Phe
65 70 75 80

Arg Ile

<210> 59
<211> 80
<212> PRT
<213> Homo sapiens

<400> 59
Lys Ala Ala Glu Glu Val Lys His Leu Lys Thr Lys Pro Ala Asp Glu
1 5 10 15
Glu Met Leu Phe Ile Tyr Ser His Tyr Lys Gln Ala Thr Val Gly Asp
20 25 30
Ile Asn Thr Glu Arg Pro Gly Met Leu Asp Phe Lys Gly Lys Ala Lys
35 40 45
Trp Asp Ala Trp Asn Glu Leu Lys Gly Thr Ser Lys Glu Asp Ala Met
50 55 60
Lys Ala Tyr Ile Asp Lys Val Glu Glu Leu Lys Lys Lys Tyr Gly Ile
65 70 75 80

<210> 60
<211> 91
<212> PRT

<213> Homo sapiens

<400> 60

Glu Lys Lys Lys Lys Lys Arg Cys Ala Gly Ile Lys His Phe Lys Thr
1 5 10 15

Lys Pro Ala Asp Asp Glu Met Arg Phe Leu Tyr Gly His Tyr Lys Arg
20 25 30

Ala Thr Val Gly Asn Ile Lys Thr Glu Arg Pro Gly Met Val Asp Phe
35 40 45

Lys Gly Lys Ala Lys Trp Asp Pro Trp Asn Leu Val Lys Gly Ala Ala
50 55 60

Arg Glu Asp Pro Met Lys Ala Lys Ala Tyr Val Lys Lys Val Glu Glu
65 70 75 80

Leu Lys Lys Lys Phe Arg Ile Arg Glu Thr Gly
85 90

<210> 61

<211> 88

<212> PRT

<213> Homo sapiens

<400> 61

Glu Ala Glu Phe Glu Lys Ala Ala Glu Glu Val Arg His Leu Lys Thr
1 5 10 15

Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His Tyr Lys Gln
20 25 30

Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu Asp Phe
35 40 45

Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys Gly Thr Ser
50 55 60

Lys Glu Asp Ala Met Lys Ala Tyr Ile Asn Lys Val Glu Glu Leu Lys
65 70 75 80

Lys Lys Tyr Gly Ile Glu Thr Gly
85

<210> 62

<211> 138

<212> PRT

<213> Homo sapiens

<400> 62

Met Ala Lys Pro Ile Ser Thr Lys Asn Thr Lys Ile Ser Arg His Gly
1 5 10 15

Trp His Ala Ala Val Ile Thr Ala Ala Arg Glu Ala Glu Ala Glu Asn

20	25	30
His Leu Ser Trp Glu Glu Lys Lys Lys Lys Lys Arg Cys Ala Gly Ile		
35	40	45
Lys His Phe Lys Thr Lys Pro Ala Asp Asp Glu Met Arg Phe Leu Tyr		
50	55	60
Gly His Tyr Lys Arg Ala Thr Val Gly Asn Ile Lys Thr Glu Arg Pro		
65	70	75
Gly Met Val Asp Phe Lys Gly Lys Ala Lys Trp Asp Pro Trp Asn Leu		
85	90	95
Val Lys Gly Ala Ala Arg Glu Asp Pro Met Lys Ala Lys Ala Tyr Val		
100	105	110
Lys Lys Val Glu Glu Leu Lys Lys Lys Phe Arg Ile Arg Glu Thr Gly		
115	120	125
Ile Val Ala Ser His Ala Phe Val Leu Asn		
130	135	

<210> 63
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 63
Ser Gln Ala Glu Phe Asp Lys Ala Ala Glu Glu Val Lys His Leu Lys
1 5 10 15
Thr Lys Pro Ala Asp Glu Glu Met Leu Phe Ile Tyr Ser His Tyr Lys
20 25 30
Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu Asp
35 40 45
Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys Gly Thr
50 55 60
Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asp Lys Val Glu Glu Leu
65 70 75 80
Lys Lys Lys Tyr Gly Ile
85

<210> 64
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 64
Ser Gln Ala Glu Phe Glu Lys Ala Ala Glu Glu Val Arg His Leu Lys
1 5 10 15

Thr Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His Tyr Lys
 20 25 30

Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu Asp
 35 40 45

Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys Gly Thr
 50 55 60

Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asn Lys Val Glu Glu Leu
 65 70 75 80

Lys Lys Lys Tyr Gly Ile
 85

<210> 65

<211> 256

<212> DNA

<213> Homo sapiens

<400> 65

aggctgattt tgacagggct gcagaagatg tgaggaagct gaaagcaaga ccagatgatg 60
 gagaactgaa agaactctat gggctttaca aacaagcaat agttggagac attaatttg 120
 cgtgtccagg aatgctagat ttaaaaggca aagccaaatg ggaagcatgg aacctcaaaa 180
 aagggttgtc gacggaagat gcgacgagtg cctatatattc taaagcaaag gagctgatag 240
 aaaaatacgg aattta 256

<210> 66

<211> 256

<212> DNA

<213> Homo sapiens

<400> 66

aggcagattt tgacaaagca gcaggggatg taaagaaatt gaaaacaaaa ccaactgacg 60
 atgaactgaa ggaactgtac ggactctaca agcagtcac tgttggggac ataatatag 120
 agtgtcctgg catgctagat ctgaagggca aggccaagtg ggacgcatgg aacctaaaga 180
 aaggcttgtc taaggaagat gcgatgagcg cttatgtttc taaagcccat gagctgatag 240
 aaaaatatgg cctgta 256

<210> 67

<211> 258

<212> DNA

<213> Homo sapiens

<400> 67

aggctgattt tgacagggct gcagaagatg tgaggaagct gaaagcaaga ccagatgatg 60
 gagaactgaa agaactctat gggctttaca aacaagcaat agttggagac attaatttg 120
 cgtgtccagg aatgctagat ttaaaaggca aagccaaatg ggaagcatgg aacctcaaaa 180
 aagggttgtc gacggaagat gcgacgagtg cctatatattc taaagcaaag gagctgatag 240
 aaaaatacgg aatttaga 258

<210> 68

<211> 259
 <212> DNA
 <213> Homo sapiens

<400> 68
 aggctgagtt tgagaaagct gcagaggagg ttaggcacct taagaccaag ccatcggatg 60
 aggagatgct gttcatctat ggccactaca aacaagcaac tgtgggacgac ataaatacag 120
 aacggcccgg gatgttggac ttcacgggca aggccaagtg ggatgcctgg aatgagctga 180
 aagggacttc caaggaagat gccatgaaag cttacatcaa caaagtagaa gagctaaaga 240
 aaaaatacgg gatatgaga 259

<210> 69
 <211> 88
 <212> PRT
 <213> Homo sapiens

<400> 69
 Phe Phe Leu Lys Ala Asp Phe Asp Arg Ala Ala Glu Asp Val Arg Lys
 1 5 10 15
 Leu Lys Ala Arg Pro Asp Asp Gly Glu Leu Lys Glu Leu Tyr Gly Leu
 20 25 30
 Tyr Lys Gln Ala Ile Val Gly Asp Ile Asn Ile Ala Cys Pro Gly Met
 35 40 45
 Leu Asp Leu Lys Gly Lys Ala Lys Trp Glu Ala Trp Asn Leu Lys Lys
 50 55 60
 Gly Leu Ser Thr Glu Asp Ala Thr Ser Ala Tyr Ile Ser Lys Ala Lys
 65 70 75 80
 Glu Leu Ile Glu Lys Tyr Gly Ile
 85

<210> 70
 <211> 89
 <212> PRT
 <213> Homo sapiens

<400> 70
 Phe Phe Leu His Gln Ala Asp Phe Asp Glu Ala Ala Glu Glu Val Lys
 1 5 10 15
 Lys Leu Lys Thr Arg Pro Thr Asp Glu Glu Leu Lys Glu Leu Tyr Gly
 20 25 30
 Phe Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Ile Glu Cys Pro Gly
 35 40 45
 Met Leu Asp Leu Lys Gly Lys Ala Lys Trp Glu Ala Trp Asn Leu Lys
 50 55 60
 Lys Gly Ile Ser Lys Glu Asp Ala Met Asn Ala Tyr Ile Ser Lys Ala
 65 70 75 80

Lys Thr Met Val Glu Lys Tyr Gly Ile
85

<210> 71
<211> 85
<212> PRT
<213> Homo sapiens

<400> 71
Lys Ala Asp Phe Asp Arg Ala Ala Glu Asp Val Arg Lys Leu Lys Ala
1 5 10 15
Arg Pro Asp Asp Gly Glu Leu Lys Glu Leu Tyr Gly Leu Tyr Lys Gln
20 25 30
Ala Ile Val Gly Asp Ile Asn Ile Ala Cys Pro Gly Met Leu Asp Leu
35 40 45
Lys Gly Lys Ala Lys Trp Glu Ala Trp Asn Leu Lys Lys Gly Leu Ser
50 55 60
Thr Glu Asp Ala Thr Ser Ala Tyr Ile Ser Lys Ala Lys Glu Leu Ile
65 70 75 80
Glu Lys Tyr Gly Ile
85

<210> 72
<211> 85
<212> PRT
<213> Homo sapiens

<220>
<221> VARIANT
<222> (1)..(85)
<223> Wherein Xaa is any amino acid.

<400> 72
Xaa Ala Asp Phe Asp Xaa Ala Ala Xaa Asp Val Xaa Lys Leu Lys Xaa
1 5 10 15
Xaa Pro Xaa Asp Xaa Glu Leu Lys Glu Leu Tyr Gly Leu Tyr Lys Gln
20 25 30
Xaa Xaa Val Gly Asp Ile Asn Ile Xaa Cys Pro Gly Met Leu Asp Leu
35 40 45
Lys Gly Lys Ala Lys Trp Xaa Ala Trp Asn Leu Lys Lys Gly Leu Ser
50 55 60
Xaa Glu Asp Ala Xaa Ser Ala Tyr Xaa Ser Lys Ala Xaa Glu Leu Ile
65 70 75 80
Glu Lys Tyr Gly Xaa

<210> 73
 <211> 85
 <212> PRT
 <213> Homo sapiens

<400> 73
 Gln Ala Asp Phe Asp Lys Ala Ala Gly Asp Val Lys Lys Leu Lys Thr
 1 5 10 15
 Lys Pro Thr Asp Asp Glu Leu Lys Glu Leu Tyr Gly Leu Tyr Lys Gln
 20 25 30
 Ser Thr Val Gly Asp Ile Asn Ile Glu Cys Pro Gly Met Leu Asp Leu
 35 40 45
 Lys Gly Lys Ala Lys Trp Asp Ala Trp Asn Leu Lys Lys Gly Leu Ser
 50 55 60
 Lys Glu Asp Ala Met Ser Ala Tyr Val Ser Lys Ala His Glu Leu Ile
 65 70 75 80
 Glu Lys Tyr Gly Leu
 85

<210> 74
 <211> 96
 <212> PRT
 <213> Homo sapiens

<400> 74
 Met Leu Leu Leu Phe Val Cys Leu Phe Phe Leu Lys Ala Asp Phe Asp
 1 5 10 15
 Arg Ala Ala Glu Asp Val Arg Lys Leu Lys Ala Arg Pro Asp Asp Gly
 20 25 30
 Glu Leu Lys Glu Leu Tyr Gly Leu Tyr Lys Gln Ala Ile Val Gly Asp
 35 40 45
 Ile Asn Ile Ala Cys Pro Gly Met Leu Asp Leu Lys Gly Lys Ala Lys
 50 55 60
 Trp Glu Ala Trp Asn Leu Lys Lys Gly Leu Ser Thr Glu Asp Ala Thr
 65 70 75 80
 Ser Ala Tyr Ile Ser Lys Ala Lys Glu Leu Ile Glu Lys Tyr Gly Ile
 85 90 95

<210> 75

<211> 88
 <212> PRT
 <213> Homo sapiens

<400> 75
 Met Ser Pro Gln Ala Asp Phe Asp Lys Ala Ala Gly Asp Val Lys Lys
 1 5 10 15
 Leu Lys Thr Lys Pro Thr Asp Asp Glu Leu Lys Glu Leu Tyr Gly Leu
 20 25 30
 Tyr Lys Gln Ser Thr Val Gly Asp Ile Asn Ile Glu Cys Pro Gly Met
 35 40 45
 Leu Asp Leu Lys Gly Lys Ala Lys Trp Asp Ala Trp Asn Leu Lys Lys
 50 55 60
 Gly Leu Ser Lys Glu Asp Ala Met Ser Ala Tyr Val Ser Lys Ala His
 65 70 75 80
 Glu Leu Ile Glu Lys Tyr Gly Leu
 85

<210> 76
 <211> 103
 <212> PRT
 <213> Homo sapiens

<400> 76
 Met Phe Gln Ala His Leu Leu Arg Gly Thr Leu Thr Leu Ser Phe Phe
 1 5 10 15
 Leu His Gln Ala Asp Phe Asp Glu Ala Ala Glu Glu Val Lys Lys Leu
 20 25 30
 Lys Thr Arg Pro Thr Asp Glu Glu Leu Lys Glu Leu Tyr Gly Phe Tyr
 35 40 45
 Lys Gln Ala Thr Val Gly Asp Ile Asn Ile Glu Cys Pro Gly Met Leu
 50 55 60
 Asp Leu Lys Gly Lys Ala Lys Trp Glu Ala Trp Asn Leu Lys Lys Gly
 65 70 75 80
 Ile Ser Lys Glu Asp Ala Met Asn Ala Tyr Ile Ser Lys Ala Lys Thr
 85 90 95
 Met Val Glu Lys Tyr Gly Ile
 100

<210> 77
 <211> 87
 <212> PRT
 <213> Homo sapiens

<400> 77

Met Ser Gln Ala Glu Phe Glu Lys Ala Ala Glu Glu Val Arg His Leu
1 5 10 15

Lys Thr Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His Tyr
20 25 30

Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu
35 40 45

Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys Gly
50 55 60

Thr Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asn Lys Val Glu Glu
65 70 75 80

Leu Lys Lys Lys Tyr Gly Ile
85

<210> 78

<211> 274

<212> DNA

<213> Homo sapiens

<400> 78

ccaccatggc actgcaggct gaattcgaca aggctgcaga agacgtgagg aagctgccaa 60
caagaccagc agataataaa gaactgaaaa aactcgatgg actttacaaa caagctataa 120
ttggagacat taatattgag tatctgggaa tgctggactt taagggcaag gccaaatgcy 180
cagcatggac cctccaaaaa aggttgtcaa aggaagatgc aacgagtgtc tctattttcta 240
aggcaaaaga gccgatagaa aaataggaca tttta 274

<210> 79

<211> 271

<212> DNA

<213> Homo sapiens

<400> 79

caaccatgtc accccaggca gatttttgaca aagcagcagg ggatgttaaag aaattgaaaa 60
caaaaccaac tgacgatgaa ctgaaggaac tgtacggact ctacaagcag tccactgttg 120
gggacataaa tatagagtgt cctggcatgc tagatctgaa gggcaaggcc aagtgggacg 180
catggaacct aaagaaaggc ttgtctaagg aagatgcgat gagcgcttat gtttctaaag 240
cccatgagct gatagaaaaa tatggcctgt a 271

<210> 80

<211> 262

<212> DNA

<213> Homo sapiens

<400> 80

caggctgaat tcgacaaggc tgcagaagac gtgaggaagc tgccaacaag accagcagat 60
aataaagaac tgaaaaaact cgatggactt tacaacaag ctataattgg agacattaat 120
attgagtatc tgggaatgct ggacttttaag ggcaaggcca aatgcgcagc atggaccctc 180
caaaaaaggt tgtcaaagga agatgcaacg agtgtctcta tttctaaggc aaaagagccg 240
atagaaaaat aggacattta ga 262

<210> 81
 <211> 260
 <212> DNA
 <213> Homo sapiens

<400> 81
 caggctgagt ttgagaaagc tgcagaggag gttaggcacc ttaagaccaa gccatcggat 60
 gaggagatgc tgttcatcta tggccactac aaacaagcaa ctgtgggcca cataaataca 120
 gaacggcccg ggatgttga cttcacgggc aaggccaagt gggatgcctg gaatgagctg 180
 aaagggactt ccaaggaaga tgccatgaaa gcttacatca acaaagtaga agagctaaag 240
 aaaaaatacg ggatatgaga 260

<210> 82
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 82
 Met Ala Leu Gln Ala Glu Phe Asp Lys Ala Ala Glu Asp Val Arg Lys
 1 5 10 15
 Leu Pro Thr Arg Pro Ala Asp Asn Lys Glu Leu Lys Lys Leu Asp Gly
 20 25 30
 Leu Tyr Lys Gln Ala Ile Ile Gly Asp Ile Asn Ile Glu Tyr Leu Gly
 35 40 45
 Met Leu Asp Phe Lys Gly Lys Ala Lys Cys Ala Ala Trp Thr Leu Gln
 50 55 60
 Lys Arg Leu Ser Lys Glu Asp Ala Thr Ser Val Ser Ile Ser Lys Ala
 65 70 75 80
 Lys Glu Pro Ile Glu Lys
 85

<210> 83
 <211> 85
 <212> PRT
 <213> Homo sapiens

<400> 83
 Met Ser Pro Gln Ala Asp Phe Asp Lys Ala Ala Gly Asp Val Lys Lys
 1 5 10 15
 Leu Lys Thr Lys Pro Thr Asp Asp Glu Leu Lys Glu Leu Tyr Gly Leu
 20 25 30
 Tyr Lys Gln Ser Thr Val Gly Asp Ile Asn Ile Glu Cys Pro Gly Met
 35 40 45
 Leu Asp Leu Lys Gly Lys Ala Lys Trp Asp Ala Trp Asn Leu Lys Lys
 50 55 60

Gly Leu Ser Lys Glu Asp Ala Met Ser Ala Tyr Val Ser Lys Ala His
65 70 75 80

Glu Leu Ile Glu Lys
85

<210> 84

<211> 88

<212> PRT

<213> Homo sapiens

<400> 84

Met Ser Pro Gln Ala Asp Phe Asp Lys Ala Ala Gly Asp Val Lys Lys
1 5 10 15

Leu Lys Thr Lys Pro Thr Asp Asp Glu Leu Lys Glu Leu Tyr Gly Leu
20 25 30

Tyr Lys Gln Ser Thr Val Gly Asp Ile Asn Ile Glu Cys Pro Gly Met
35 40 45

Leu Asp Leu Lys Gly Lys Ala Lys Trp Asp Ala Trp Asn Leu Lys Lys
50 55 60

Gly Leu Ser Lys Glu Asp Ala Met Ser Ala Tyr Val Ser Lys Ala His
65 70 75 80

Glu Leu Ile Glu Lys Tyr Gly Leu
85

<210> 85

<211> 103

<212> PRT

<213> Homo sapiens

<400> 85

Met Phe Gln Ala His Leu Leu Arg Gly Thr Leu Thr Leu Ser Phe Phe
1 5 10 15

Leu His Gln Ala Asp Phe Asp Glu Ala Ala Glu Glu Val Lys Lys Leu
20 25 30

Lys Thr Arg Pro Thr Asp Glu Glu Leu Lys Glu Leu Tyr Gly Phe Tyr
35 40 45

Lys Gln Ala Thr Val Gly Asp Ile Asn Ile Glu Cys Pro Gly Met Leu
50 55 60

Asp Leu Lys Gly Lys Ala Lys Trp Glu Ala Trp Asn Leu Lys Lys Gly
65 70 75 80

Ile Ser Lys Glu Asp Ala Met Asn Ala Tyr Ile Ser Lys Ala Lys Thr
85 90 95

Met Val Glu Lys Tyr Gly Ile
100

<210> 86
<211> 87
<212> PRT
<213> Homo sapiens

<400> 86
Met Ser Gln Ala Glu Phe Glu Lys Ala Ala Glu Glu Val Arg His Leu
1 5 10 15
Lys Thr Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His Tyr
20 25 30
Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu
35 40 45
Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys Gly
50 55 60
Thr Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asn Lys Val Glu Glu
65 70 75 80
Leu Lys Lys Lys Tyr Gly Ile
85

<210> 87
<211> 86
<212> PRT
<213> Homo sapiens

<400> 87
Met Ala Leu Gln Ala Glu Phe Asp Lys Ala Ala Glu Asp Val Arg Lys
1 5 10 15
Leu Pro Thr Arg Pro Ala Asp Asn Lys Glu Leu Lys Lys Leu Asp Gly
20 25 30
Leu Tyr Lys Gln Ala Ile Ile Gly Asp Ile Asn Ile Glu Tyr Leu Gly
35 40 45
Met Leu Asp Phe Lys Gly Lys Ala Lys Cys Ala Ala Trp Thr Leu Gln
50 55 60
Lys Arg Leu Ser Lys Glu Asp Ala Thr Ser Val Ser Ile Ser Lys Ala
65 70 75 80
Lys Glu Pro Ile Glu Lys
85

<210> 88
<211> 530
<212> PRT

<213> Homo sapiens

<400> 88

Met	Phe	Gln	Phe	His	Ala	Gly	Ser	Trp	Glu	Ser	Trp	Cys	Cys	Cys	Cys	
1				5					10					15		
Leu	Ile	Pro	Ala	Asp	Arg	Pro	Trp	Asp	Arg	Gly	Gln	His	Trp	Gln	Leu	
			20					25					30			
Glu	Met	Ala	Asp	Thr	Arg	Ser	Val	His	Glu	Thr	Arg	Phe	Glu	Ala	Ala	
		35					40					45				
Val	Lys	Val	Ile	Gln	Ser	Leu	Pro	Lys	Asn	Gly	Ser	Phe	Gln	Pro	Thr	
	50					55					60					
Asn	Glu	Met	Met	Leu	Lys	Phe	Tyr	Ser	Phe	Tyr	Lys	Gln	Ala	Thr	Glu	
65					70					75					80	
Gly	Pro	Cys	Lys	Leu	Ser	Arg	Pro	Gly	Phe	Trp	Asp	Pro	Ile	Gly	Arg	
				85					90						95	
Tyr	Lys	Trp	Asp	Ala	Trp	Ser	Ser	Leu	Gly	Asp	Met	Thr	Lys	Glu	Glu	
			100					105					110			
Ala	Met	Ile	Ala	Tyr	Val	Glu	Glu	Met	Lys	Lys	Ile	Ile	Glu	Thr	Met	
		115						120					125			
Pro	Met	Thr	Glu	Lys	Val	Glu	Glu	Leu	Leu	Arg	Val	Ile	Gly	Pro	Phe	
	130							135				140				
Tyr	Glu	Ile	Val	Glu	Asp	Lys	Lys	Ser	Gly	Arg	Ser	Ser	Asp	Ile	Thr	
145					150					155					160	
Ser	Val	Arg	Leu	Glu	Lys	Ile	Ser	Lys	Cys	Leu	Glu	Asp	Leu	Gly	Asn	
			165						170					175		
Val	Leu	Thr	Ser	Thr	Pro	Asn	Ala	Lys	Thr	Val	Asn	Gly	Lys	Ala	Glu	
			180						185				190			
Ser	Ser	Asp	Ser	Gly	Ala	Glu	Ser	Glu	Glu	Glu	Glu	Ala	Gln	Glu	Glu	
		195						200					205			
Val	Lys	Gly	Ala	Glu	His	Ser	Asp	Asn	Asp	Lys	Lys	Met	Met	Lys	Lys	
	210						215					220				
Ser	Ala	Asp	His	Lys	Asn	Leu	Glu	Val	Ile	Val	Thr	Asn	Gly	Tyr	Asp	
225					230					235					240	
Lys	Asp	Gly	Phe	Val	Gln	Asp	Ile	Gln	Asn	Asp	Ile	His	Ala	Ser	Ser	
			245						250					255		
Ser	Leu	Asn	Gly	Arg	Ser	Thr	Glu	Glu	Val	Lys	Pro	Ile	Asp	Glu	Asn	
			260					265					270			
Leu	Gly	Gln	Thr	Gly	Lys	Ser	Ala	Val	Cys	Ile	His	Gln	Gly	Ile	Asn	
		275					280						285			

Asp Asp His Val Glu Asp Val Thr Gly Ile Gln His Leu Thr Ser Asp
 290 295 300
 Ser Asp Ser Glu Val Tyr Cys Asp Ser Met Glu Gln Phe Gly Gln Glu
 305 310 315 320
 Glu Ser Leu Asp Ser Phe Thr Ser Asn Asn Gly Pro Phe Gln Tyr Tyr
 325 330 335
 Leu Gly Gly His Ser Ser Gln Pro Met Glu Asn Ser Gly Phe Arg Glu
 340 345 350
 Asp Ile Gln Val Pro Pro Gly Asn Gly Asn Ile Gly Asn Met Gln Val
 355 360 365
 Val Ala Val Glu Gly Lys Gly Glu Val Lys His Gly Gly Glu Asp Gly
 370 375 380
 Arg Asn Asn Ser Gly Ala Pro His Arg Glu Lys Arg Gly Gly Glu Thr
 385 390 395 400
 Asp Glu Phe Ser Asn Val Arg Arg Gly Arg Gly His Arg Met Gln His
 405 410 415
 Leu Ser Glu Gly Thr Lys Gly Arg Gln Val Gly Ser Gly Gly Asp Gly
 420 425 430
 Glu Arg Trp Gly Ser Asp Arg Gly Ser Arg Gly Ser Leu Asn Glu Gln
 435 440 445
 Ile Ala Leu Val Leu Met Arg Leu Gln Glu Asp Met Gln Asn Val Leu
 450 455 460
 Gln Arg Leu Gln Lys Leu Glu Thr Leu Thr Ala Ala Lys Ser Ser Thr
 465 470 475 480
 Ser Thr Leu Gln Thr Ala Pro Gln Pro Thr Ser Ser Gln Arg Pro Ser
 485 490 495
 Trp Trp Pro Phe Glu Met Ser Pro Gly Val Leu Thr Phe Ala Ile Ile
 500 505 510
 Trp Pro Phe Ile Ala Gln Trp Leu Val Tyr Leu Tyr Tyr Gln Arg Arg
 515 520 525
 Arg Arg
 530

<210> 89

<211> 530

<212> PRT

<213> Homo sapiens

<400> 89

Met Phe Gln Phe His Ala Gly Ser Trp Glu Ser Trp Cys Cys Cys Cys
 1 5 10 15

Cys Leu Ile Pro Gly Asp Arg Pro Trp Asp Arg Gly Arg Arg Trp Arg
 20 25 30
 Leu Glu Met Arg His Thr Arg Ser Val His Glu Thr Arg Phe Glu Ala
 35 40 45
 Ala Val Lys Val Ile Gln Ser Leu Pro Lys Asn Gly Ser Phe Gln Pro
 50 55 60
 Thr Asn Glu Met Met Leu Lys Phe Tyr Ser Phe Tyr Lys Gln Ala Thr
 65 70 75 80
 Glu Gly Pro Cys Lys Leu Ser Lys Pro Gly Phe Trp Asp Pro Val Gly
 85 90 95
 Arg Tyr Lys Trp Asp Ala Trp Ser Ser Leu Gly Asp Met Thr Lys Glu
 100 105 110
 Glu Ala Met Ile Ala Tyr Val Glu Glu Met Lys Lys Ile Leu Glu Thr
 115 120 125
 Met Pro Met Thr Glu Lys Val Glu Glu Leu Leu His Val Ile Gly Pro
 130 135 140
 Phe Tyr Glu Ile Val Glu Asp Lys Lys Ser Gly Arg Ser Ser Asp Leu
 145 150 155 160
 Thr Ser Val Arg Leu Glu Lys Ile Ser Lys Cys Leu Glu Asp Leu Gly
 165 170 175
 Asn Val Leu Ala Ser Thr Pro Asn Ala Lys Thr Val Asn Gly Lys Ala
 180 185 190
 Glu Ser Ser Asp Ser Gly Ala Glu Ser Glu Glu Glu Ala Ala Gln Glu
 195 200 205
 Asp Pro Lys Arg Pro Glu Pro Arg Asp Ser Asp Lys Lys Met Met Lys
 210 215 220
 Lys Ser Ala Asp His Lys Asn Leu Glu Ile Ile Val Thr Asn Gly Tyr
 225 230 235 240
 Asp Lys Asp Ser Phe Val Gln Gly Val Gln Asn Ser Ile His Thr Ser
 245 250 255
 Pro Ser Leu Asn Gly Arg Cys Thr Glu Glu Val Lys Ser Val Asp Glu
 260 265 270
 Asn Leu Glu Gln Thr Gly Lys Thr Val Val Phe Val His Gln Asp Val
 275 280 285
 Asn Ser Asp His Val Glu Asp Ile Ser Gly Ile Gln His Leu Thr Ser
 290 295 300
 Asp Ser Asp Ser Glu Val Tyr Cys Asp Ser Met Glu Gln Phe Gly Gln
 305 310 315 320

Glu Glu Ser Leu Asp Gly Phe Ile Ser Asn Asn Gly Pro Phe Ser Tyr
 325 330 335
 Tyr Leu Gly Gly Asn Pro Ser Gln Pro Leu Glu Ser Ser Gly Phe Pro
 340 345 350
 Glu Ala Val Gln Gly Leu Pro Gly Asn Gly Ser Pro Glu Asp Met Gln
 355 360 365
 Gly Ala Val Val Glu Gly Lys Gly Glu Val Lys Arg Gly Gly Glu Asp
 370 375 380
 Gly Gly Ser Asn Ser Gly Ala Pro His Arg Glu Lys Arg Ala Gly Glu
 385 390 395 400
 Ser Glu Glu Phe Ser Asn Ile Arg Arg Gly Arg Gly His Arg Met Gln
 405 410 415
 His Leu Ser Glu Gly Ser Lys Gly Arg Gln Val Gly Ser Gly Gly Asp
 420 425 430
 Gly Glu Arg Trp Gly Ser Asp Arg Gly Ser Arg Gly Ser Leu Asn Glu
 435 440 445
 Gln Ile Ala Leu Val Leu Met Arg Leu Gln Glu Asp Met Gln Asn Val
 450 455 460
 Leu Gln Arg Leu His Lys Leu Glu Met Leu Ala Ala Ser Gln Ala Lys
 465 470 475 480
 Ser Ser Ala Leu Gln Thr Ser Asn Gln Pro Thr Ser Pro Arg Pro Ser
 485 490 495
 Trp Trp Pro Phe Glu Met Ser Pro Gly Ala Leu Thr Phe Ala Ile Ile
 500 505 510
 Trp Pro Phe Ile Ala Gln Trp Leu Val His Leu Tyr Tyr Gln Arg Arg
 515 520 525
 Arg Arg
 530

<210> 90
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 90
 Met Ser Gln Ala Phe Glu Lys Ala Ala Lys Asp Ile Lys His Leu Glu
 1 5 10 15
 Thr Lys Pro Ala Asp Asp Glu Arg Met Phe Ile Tyr Ser Arg Cys Lys
 20 25 30
 Gln Ala Thr Val His Asp Leu Asn Thr Glu Trp Pro Arg Met Leu Asp

35 40 45
 Leu Lys Gly Lys Ala Lys Gln Asp Ala Trp Asn Glu Leu Lys Asp Thr
 50 55 60
 Ala Lys Glu Asp Ala Val Lys Ala Asp Ile Asn Lys Val Glu Glu Arg
 65 70 75 80
 Asn Lys Lys Tyr Arg Ile
 85

<210> 91
 <211> 87
 <212> PRT
 <213> Homo sapiens

<400> 91
 Met Ser Gln Ala Glu Phe Asp Lys Ala Ala Glu Glu Val Lys His Leu
 1 5 10 15
 Lys Thr Lys Pro Ala Asp Glu Glu Met Leu Phe Ile Tyr Ser His Tyr
 20 25 30
 Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu
 35 40 45
 Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys Gly
 50 55 60
 Thr Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asp Lys Val Glu Glu
 65 70 75 80
 Leu Lys Lys Lys Tyr Gly Ile
 85

<210> 92
 <211> 104
 <212> PRT
 <213> Homo sapiens

<400> 92
 Met Trp Gly Asp Leu Trp Leu Leu Pro Pro Ala Ser Ala Asn Pro Gly
 1 5 10 15
 Thr Gly Thr Glu Ala Glu Phe Glu Lys Ala Ala Glu Glu Val Arg His
 20 25 30
 Leu Lys Thr Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His
 35 40 45
 Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met
 50 55 60
 Leu Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys
 65 70 75 80

Gly Thr Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asn Lys Val Glu
85 90 95

Glu Leu Lys Lys Lys Tyr Gly Ile
100

<210> 93
<211> 104
<212> PRT
<213> Homo sapiens

<400> 93
Met Trp Gly Asp Leu Trp Leu Leu Pro Pro Ala Ser Ala Asn Pro Gly
1 5 10 15

Thr Gly Thr Glu Ala Glu Phe Glu Lys Ala Ala Glu Glu Val Arg His
20 25 30

Leu Lys Thr Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His
35 40 45

Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met
50 55 60

Leu Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys
65 70 75 80

Gly Thr Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asn Lys Val Glu
85 90 95

Glu Leu Lys Lys Lys Tyr Gly Ile
100

<210> 94
<211> 359
<212> PRT
<213> Homo sapiens

<400> 94
Met Arg Ala Ser Gln Lys Asp Phe Glu Asn Ser Met Asn Gln Val Lys
1 5 10 15

Leu Leu Lys Lys Asp Pro Gly Asn Glu Val Lys Leu Lys Leu Tyr Ala
20 25 30

Leu Tyr Lys Gln Ala Thr Glu Gly Pro Cys Asn Met Pro Lys Pro Gly
35 40 45

Val Phe Asp Leu Ile Asn Lys Ala Lys Trp Asp Ala Trp Asn Ala Leu
50 55 60

Gly Ser Leu Pro Lys Glu Ala Ala Arg Gln Asn Tyr Val Asp Leu Val
65 70 75 80

Ser Ser Leu Ser Pro Ser Leu Glu Ser Ser Ser Gln Val Glu Pro Gly
 85 90 95
 Thr Asp Arg Lys Ser Thr Gly Phe Glu Thr Leu Val Val Thr Ser Glu
 100 105 110
 Asp Gly Ile Thr Lys Ile Met Phe Asn Arg Pro Lys Lys Lys Asn Ala
 115 120 125
 Ile Asn Thr Glu Met Tyr His Glu Ile Met Arg Ala Leu Lys Ala Ala
 130 135 140
 Ser Lys Asp Asp Ser Ile Ile Thr Val Leu Thr Gly Asn Gly Asp Tyr
 145 150 155 160
 Tyr Ser Ser Gly Asn Asp Leu Thr Asn Phe Thr Asp Ile Pro Pro Gly
 165 170 175
 Gly Val Glu Glu Lys Ala Lys Asn Asn Ala Val Leu Leu Arg Glu Phe
 180 185 190
 Val Gly Cys Phe Ile Asp Phe Pro Lys Pro Leu Ile Ala Val Val Asn
 195 200 205
 Gly Pro Ala Val Gly Ile Ser Val Thr Leu Leu Gly Leu Phe Asp Ala
 210 215 220
 Val Tyr Ala Ser Asp Arg Ala Thr Phe His Thr Pro Phe Ser His Leu
 225 230 235 240
 Gly Gln Ser Pro Glu Gly Cys Ser Ser Tyr Thr Phe Pro Lys Ile Met
 245 250 255
 Ser Pro Ala Lys Ala Thr Glu Met Leu Ile Phe Gly Lys Lys Leu Thr
 260 265 270
 Ala Gly Glu Ala Cys Ala Gln Gly Leu Val Thr Glu Val Phe Pro Asp
 275 280 285
 Ser Thr Phe Gln Lys Glu Val Trp Thr Arg Leu Lys Ala Phe Ala Lys
 290 295 300
 Leu Pro Pro Asn Ala Leu Arg Ile Ser Lys Glu Val Ile Arg Lys Arg
 305 310 315 320
 Glu Arg Glu Lys Leu His Ala Val Asn Ala Glu Glu Cys Asn Val Leu
 325 330 335
 Gln Gly Arg Trp Leu Ser Asp Glu Cys Thr Asn Ala Val Val Asn Phe
 340 345 350
 Leu Ser Arg Lys Ser Lys Leu
 355

<210> 95
 <211> 359

<212> PRT

<213> Homo sapiens

<400> 95

Met	Arg	Ala	Ser	Gln	Lys	Asp	Phe	Glu	Asn	Ser	Met	Asn	Gln	Val	Lys	
1				5					10					15		
Leu	Leu	Lys	Lys	Asp	Pro	Gly	Asn	Glu	Val	Lys	Leu	Lys	Leu	Tyr	Ala	
			20					25					30			
Leu	Tyr	Lys	Gln	Ala	Thr	Glu	Gly	Pro	Cys	Asn	Met	Pro	Lys	Pro	Gly	
		35					40					45				
Val	Phe	Asp	Leu	Ile	Asn	Lys	Ala	Lys	Trp	Asp	Ala	Trp	Asn	Ala	Leu	
	50					55					60					
Gly	Ser	Leu	Pro	Lys	Glu	Ala	Ala	Arg	Gln	Asn	Tyr	Val	Asp	Leu	Val	
65					70					75					80	
Ser	Ser	Leu	Ser	Pro	Ser	Leu	Glu	Ser	Ser	Ser	Gln	Val	Glu	Pro	Gly	
				85					90					95		
Thr	Asp	Arg	Lys	Ser	Thr	Gly	Phe	Glu	Thr	Leu	Val	Val	Thr	Ser	Glu	
			100					105					110			
Asp	Gly	Ile	Thr	Lys	Ile	Met	Phe	Asn	Arg	Pro	Lys	Lys	Lys	Asn	Ala	
		115					120					125				
Ile	Asn	Thr	Glu	Met	Tyr	His	Glu	Ile	Met	Arg	Ala	Leu	Lys	Ala	Ala	
	130					135					140					
Ser	Lys	Asp	Asp	Ser	Ile	Ile	Thr	Val	Leu	Thr	Gly	Asn	Gly	Asp	Tyr	
145					150					155					160	
Tyr	Ser	Ser	Gly	Asn	Asp	Leu	Thr	Asn	Phe	Thr	Asp	Ile	Pro	Pro	Gly	
				165					170					175		
Gly	Val	Glu	Glu	Lys	Ala	Lys	Asn	Asn	Ala	Val	Leu	Leu	Arg	Glu	Phe	
			180					185					190			
Val	Gly	Cys	Phe	Ile	Asp	Phe	Pro	Lys	Pro	Leu	Ile	Ala	Val	Val	Asn	
		195					200					205				
Gly	Pro	Ala	Val	Gly	Ile	Ser	Val	Thr	Leu	Leu	Gly	Leu	Phe	Asp	Ala	
	210					215					220					
Val	Tyr	Ala	Ser	Asp	Arg	Ala	Thr	Phe	His	Thr	Pro	Phe	Ser	His	Leu	
225					230					235					240	
Gly	Gln	Ser	Pro	Glu	Gly	Cys	Ser	Ser	Tyr	Thr	Phe	Pro	Lys	Ile	Met	
				245					250					255		
Ser	Pro	Ala	Lys	Ala	Thr	Glu	Met	Leu	Ile	Phe	Gly	Lys	Lys	Leu	Thr	
			260					265					270			
Ala	Gly	Glu	Ala	Cys	Ala	Gln	Gly	Leu	Val	Thr	Glu	Val	Phe	Pro	Asp	
		275					280					285				

Ser Thr Phe Gln Lys Glu Val Trp Thr Arg Leu Lys Ala Phe Ala Lys
 290 295 300
 Leu Pro Pro Asn Ala Leu Arg Ile Ser Lys Glu Val Ile Arg Lys Arg
 305 310 315 320
 Glu Arg Glu Lys Leu His Ala Val Asn Ala Glu Glu Cys Asn Val Leu
 325 330 335
 Gln Gly Arg Trp Leu Ser Asp Glu Cys Thr Asn Ala Val Val Asn Phe
 340 345 350
 Leu Ser Arg Lys Ser Lys Leu
 355

<210> 96
 <211> 282
 <212> PRT
 <213> Homo sapiens

<400> 96
 Met Ala Ser Ser Phe Leu Pro Ala Gly Ala Ile Thr Gly Asp Ser Gly
 1 5 10 15
 Gly Glu Leu Ser Ser Gly Asp Asp Ser Gly Glu Val Glu Phe Pro His
 20 25 30
 Ser Pro Glu Ile Glu Glu Thr Ser Cys Leu Ala Glu Leu Phe Glu Lys
 35 40 45
 Ala Ala Ala His Leu Gln Gly Leu Ile Gln Val Ala Ser Arg Glu Gln
 50 55 60
 Leu Leu Tyr Leu Tyr Ala Arg Tyr Lys Gln Val Lys Val Gly Asn Cys
 65 70 75 80
 Asn Thr Pro Lys Pro Ser Phe Phe Asp Phe Glu Gly Lys Gln Lys Trp
 85 90 95
 Glu Ala Trp Lys Ala Leu Gly Asp Ser Ser Pro Ser Gln Ala Met Gln
 100 105 110
 Glu Tyr Ile Ala Val Val Lys Lys Leu Asp Pro Gly Trp Asn Pro Gln
 115 120 125
 Ile Pro Glu Lys Lys Gly Lys Glu Ala Asn Thr Gly Phe Gly Gly Pro
 130 135 140
 Val Ile Ser Ser Leu Tyr His Glu Glu Thr Ile Arg Glu Glu Asp Lys
 145 150 155 160
 Asn Ile Phe Asp Tyr Cys Arg Glu Asn Asn Ile Asp His Ile Thr Lys
 165 170 175
 Ala Ile Lys Ser Lys Asn Val Asp Val Asn Val Lys Asp Glu Glu Gly

180	185	190
Arg Ala Leu Leu His Trp Ala Cys Asp Arg Gly His Lys Glu Leu Val		
195	200	205
Thr Val Leu Leu Gln His Arg Ala Asp Ile Asn Cys Gln Asp Asn Glu		
210	215	220
Gly Gln Thr Ala Leu His Tyr Ala Ser Ala Cys Glu Phe Leu Asp Ile		
225	230	235
Val Glu Leu Leu Leu Gln Ser Gly Ala Asp Pro Thr Leu Arg Asp Gln		
245	250	255
Asp Gly Cys Leu Pro Glu Glu Val Thr Gly Cys Lys Thr Val Ser Leu		
260	265	270
Val Leu Gln Arg His Thr Thr Gly Lys Ala		
275	280	

<210> 97
 <211> 279
 <212> PRT
 <213> Homo sapiens

<400> 97
Met Ala Ser Ser Phe Leu Pro Ala Gly Ala Ile Thr Gly Asp Ser Gly
1 5 10 15
Gly Glu Leu Ser Ser Gly Asp Asp Ser Gly Glu Val Glu Phe Pro His
20 25 30
Ser Pro Glu Ile Glu Glu Thr Ser Cys Leu Ala Glu Leu Phe Glu Lys
35 40 45
Ala Ala Ala His Leu Gln Gly Leu Ile Gln Val Ala Ser Arg Glu Gln
50 55 60
Leu Leu Tyr Leu Tyr Ala Arg Tyr Lys Gln Val Lys Val Gly Asn Cys
65 70 75 80
Asn Thr Pro Lys Pro Ser Phe Phe Asp Phe Glu Gly Lys Gln Lys Trp
85 90 95
Glu Ala Trp Lys Ala Leu Gly Asp Ser Ser Pro Ser Gln Ala Met Gln
100 105 110
Glu Tyr Ile Ala Val Val Lys Lys Leu Asp Pro Gly Trp Asn Pro Gln
115 120 125
Ile Pro Glu Lys Lys Arg Lys Arg Ser Lys Tyr Lys Val Trp Ala Ser
130 135 140
Tyr Phe Ser Ile Ser Arg Asn His Gln Gly Arg Asp Lys Asn Ile Phe
145 150 155 160

Asp Tyr Cys Arg Glu Asn Asn Ile Asp His Ile Thr Lys Ala Ile Lys
 165 170 175
 Ser Lys Asn Val Asp Val Asn Val Lys Asp Glu Glu Gly Arg Ala Leu
 180 185 190
 Leu His Trp Ala Cys Asp Arg Gly His Lys Glu Leu Val Thr Val Leu
 195 200 205
 Leu Gln His Arg Ala Asp Ile Asn Cys Gln Asp Asn Glu Gly Gln Thr
 210 215 220
 Ala Leu His Tyr Ala Ser Ala Cys Glu Phe Leu Asp Ile Val Glu Leu
 225 230 235 240
 Leu Leu Gln Ser Gly Ala Asp Pro Thr Leu Arg Asp Gln Asp Gly Cys
 245 250 255
 Leu Pro Glu Glu Val Thr Gly Cys Lys Thr Val Ser Leu Val Leu Gln
 260 265 270
 Arg His Thr Thr Gly Lys Ala
 275

<210> 98
 <211> 89
 <212> PRT
 <213> Homo sapiens

<400> 98
 Thr Ala Ser Thr Thr Pro Cys Ala Lys Trp Ser Ser Ser Cys Ala Ala
 1 5 10 15
 Leu Lys Gln Leu Lys Gly Pro Val Ser Asp Gln Glu Lys Leu Leu Val
 20 25 30
 Tyr Gly Leu Tyr Lys Gln Ala Thr Gln Gly Asp Cys Asp Ile Pro Gly
 35 40 45
 Pro Pro Ala Ser Asp Val Arg Ala Arg Ala Lys Trp Glu Ala Trp Ser
 50 55 60
 Ala Asn Lys Gly Ala Ser Lys Met Asp Ala Met Arg Gly Tyr Ala Ala
 65 70 75 80
 Lys Val Glu Glu Leu Thr Lys Lys Glu
 85

<210> 99
 <211> 104
 <212> PRT
 <213> Homo sapiens

<400> 99
 Met Trp Gly Asp Leu Trp Leu Leu Pro Pro Ala Ser Ala Asn Pro Gly

1 5 10 15
 Thr Gly Thr Glu Ala Glu Phe Glu Lys Ala Ala Glu Glu Val Arg His
 20 25 30
 Leu Lys Thr Lys Pro Ser Asp Glu Glu Met Leu Phe Ile Tyr Gly His
 35 40 45
 Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met
 50 55 60
 Leu Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp Asn Glu Leu Lys
 65 70 75 80
 Gly Thr Ser Lys Glu Asp Ala Met Lys Ala Tyr Ile Asn Lys Val Glu
 85 90 95
 Glu Leu Lys Lys Lys Tyr Gly Ile
 100

<210> 100
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 100
 Met Ser Gln Ala Phe Glu Lys Ala Ala Lys Asp Ile Lys His Leu Glu
 1 5 10 15
 Thr Lys Pro Ala Asp Asp Glu Arg Met Phe Ile Tyr Ser Arg Cys Lys
 20 25 30
 Gln Ala Thr Val His Asp Leu Asn Thr Glu Trp Pro Arg Met Leu Asp
 35 40 45
 Leu Lys Gly Lys Ala Lys Gln Asp Ala Trp Asn Glu Leu Lys Asp Thr
 50 55 60
 Ala Lys Glu Asp Ala Val Lys Ala Asp Ile Asn Lys Val Glu Glu Arg
 65 70 75 80
 Asn Lys Lys Tyr Arg Ile
 85

<210> 101
 <211> 138
 <212> PRT
 <213> Homo sapiens

<400> 101
 Met Ala Lys Pro Ile Ser Thr Lys Asn Thr Lys Ile Ser Arg His Gly
 1 5 10 15
 Trp His Ala Ala Val Ile Thr Ala Ala Arg Glu Ala Glu Ala Glu Asn
 20 25 30

His Leu Ser Trp Glu Glu Lys Lys Lys Lys Lys Arg Cys Ala Gly Ile
 35 40 45
 Lys His Phe Lys Thr Lys Pro Ala Asp Asp Glu Met Arg Phe Leu Tyr
 50 55 60
 Gly His Tyr Lys Arg Ala Thr Val Gly Asn Ile Lys Thr Glu Arg Pro
 65 70 75 80
 Gly Met Val Asp Phe Lys Gly Lys Ala Lys Trp Asp Pro Trp Asn Leu
 85 90 95
 Val Lys Gly Ala Ala Arg Glu Asp Pro Met Lys Ala Lys Ala Tyr Val
 100 105 110
 Lys Lys Val Glu Glu Leu Lys Lys Lys Phe Arg Ile Arg Glu Thr Gly
 115 120 125
 Ile Val Ala Ser His Ala Phe Val Leu Asn
 130 135

<210> 102
 <211> 96
 <212> PRT
 <213> Homo sapiens

<400> 102
 Met Leu Leu Leu Phe Val Cys Leu Phe Phe Leu Lys Ala Asp Phe Asp
 1 5 10 15
 Arg Ala Ala Glu Asp Val Arg Lys Leu Lys Ala Arg Pro Asp Asp Gly
 20 25 30
 Glu Leu Lys Glu Leu Tyr Gly Leu Tyr Lys Gln Ala Ile Val Gly Asp
 35 40 45
 Ile Asn Ile Ala Cys Pro Gly Met Leu Asp Leu Lys Gly Lys Ala Lys
 50 55 60
 Trp Glu Ala Trp Asn Leu Lys Lys Gly Leu Ser Thr Glu Asp Ala Thr
 65 70 75 80
 Ser Ala Tyr Ile Ser Lys Ala Lys Glu Leu Ile Glu Lys Tyr Gly Ile
 85 90 95

<210> 103
 <211> 88
 <212> PRT
 <213> Homo sapiens

<400> 103

Met Ser Leu Gln Ala Asp Phe Asp Met Val Thr Glu Asp Val Arg Lys
1 5 10 15
Leu Lys Thr Arg Pro Asp Asp Glu Glu Leu Lys Glu Leu Tyr Gly Leu
20 25 30
Tyr Lys Gln Ala Val Ile Gly Asn Ile Asn Ile Glu Cys Ser Glu Met
35 40 45
Leu Glu Leu Lys Gly Lys Ala Lys Trp Glu Ala Gln Asn Pro Gln Lys
50 55 60
Gly Leu Ser Glu Glu Asp Met Met Arg Ala Phe Ile Ser Lys Ala Glu
65 70 75 80
Glu Leu Ile Glu Lys Tyr Gly Ile
85

<210> 104
<211> 86
<212> PRT
<213> Homo sapiens

<400> 104
Met Ala Leu Gln Ala Glu Phe Asp Lys Ala Ala Glu Asp Val Arg Lys
1 5 10 15
Leu Pro Thr Arg Pro Ala Asp Asn Lys Glu Leu Lys Lys Leu Asp Gly
20 25 30
Leu Tyr Lys Gln Ala Ile Ile Gly Asp Ile Asn Ile Glu Tyr Leu Gly
35 40 45
Met Leu Asp Phe Lys Gly Lys Ala Lys Cys Ala Ala Trp Thr Leu Gln
50 55 60
Lys Arg Leu Ser Lys Glu Asp Ala Thr Ser Val Ser Ile Ser Lys Ala
65 70 75 80
Lys Glu Pro Ile Glu Lys
85

<210> 105
<211> 282
<212> PRT
<213> Homo sapiens

<400> 105
Met Ala Ser Ser Phe Leu Pro Ala Gly Ala Ile Thr Gly Asp Ser Gly
1 5 10 15
Gly Glu Leu Ser Ser Gly Asp Asp Ser Gly Glu Val Glu Phe Pro His
20 25 30
Ser Pro Glu Ile Glu Glu Thr Ser Cys Leu Ala Glu Leu Phe Glu Lys

35	40	45
Ala Ala Ala His Leu Gln Gly Leu Ile Gln Val Ala Ser Arg Glu Gln		
50	55	60
Leu Leu Tyr Leu Tyr Ala Arg Tyr Lys Gln Val Lys Val Gly Asn Cys		
65	70	75
Asn Thr Pro Lys Pro Ser Phe Phe Asp Phe Glu Gly Lys Gln Lys Trp		
	85	90
Glu Ala Trp Lys Ala Leu Gly Asp Ser Ser Pro Ser Gln Ala Met Gln		
100	105	110
Glu Tyr Ile Ala Val Val Lys Lys Leu Asp Pro Gly Trp Asn Pro Gln		
115	120	125
Ile Pro Glu Lys Lys Gly Lys Glu Ala Asn Thr Gly Phe Gly Gly Pro		
130	135	140
Val Ile Ser Ser Leu Tyr His Glu Glu Thr Ile Arg Glu Glu Asp Lys		
145	150	155
Asn Ile Phe Asp Tyr Cys Arg Glu Asn Asn Ile Asp His Ile Thr Lys		
	165	170
Ala Ile Lys Ser Lys Asn Val Asp Val Asn Val Lys Asp Glu Glu Gly		
180	185	190
Arg Ala Leu Leu His Trp Ala Cys Asp Arg Gly His Lys Glu Leu Val		
195	200	205
Thr Val Leu Leu Gln His Arg Ala Asp Ile Asn Cys Gln Asp Asn Glu		
210	215	220
Gly Gln Thr Ala Leu His Tyr Ala Ser Ala Cys Glu Phe Leu Asp Ile		
225	230	235
Val Glu Leu Leu Leu Gln Ser Gly Ala Asp Pro Thr Leu Arg Asp Gln		
	245	250
Asp Gly Cys Leu Pro Glu Glu Val Thr Gly Cys Lys Thr Val Ser Leu		
260	265	270
Val Leu Gln Arg His Thr Thr Gly Lys Ala		
275	280	

<210> 106

<211> 359

<212> PRT

<213> Homo sapiens

<400> 106

Met Arg Ala Ser Gln Lys Asp Phe Glu Asn Ser Met Asn Gln Val Lys
1 5 10 15

Leu Leu Lys Lys Asp Pro Gly Asn Glu Val Lys Leu Lys Leu Tyr Ala
 20 25 30
 Leu Tyr Lys Gln Ala Thr Glu Gly Pro Cys Asn Met Pro Lys Pro Gly
 35 40 45
 Val Phe Asp Leu Ile Asn Lys Ala Lys Trp Asp Ala Trp Asn Ala Leu
 50 55 60
 Gly Ser Leu Pro Lys Glu Ala Ala Arg Gln Asn Tyr Val Asp Leu Val
 65 70 75 80
 Ser Ser Leu Ser Pro Ser Leu Glu Ser Ser Ser Gln Val Glu Pro Gly
 85 90 95
 Thr Asp Arg Lys Ser Thr Gly Phe Glu Thr Leu Val Val Thr Ser Glu
 100 105 110
 Asp Gly Ile Thr Lys Ile Met Phe Asn Arg Pro Lys Lys Lys Asn Ala
 115 120 125
 Ile Asn Thr Glu Met Tyr His Glu Ile Met Arg Ala Leu Lys Ala Ala
 130 135 140
 Ser Lys Asp Asp Ser Ile Ile Thr Val Leu Thr Gly Asn Gly Asp Tyr
 145 150 155 160
 Tyr Ser Ser Gly Asn Asp Leu Thr Asn Phe Thr Asp Ile Pro Pro Gly
 165 170 175
 Gly Val Glu Glu Lys Ala Lys Asn Asn Ala Val Leu Leu Arg Glu Phe
 180 185 190
 Val Gly Cys Phe Ile Asp Phe Pro Lys Pro Leu Ile Ala Val Val Asn
 195 200 205
 Gly Pro Ala Val Gly Ile Ser Val Thr Leu Leu Gly Leu Phe Asp Ala
 210 215 220
 Val Tyr Ala Ser Asp Arg Ala Thr Phe His Thr Pro Phe Ser His Leu
 225 230 235 240
 Gly Gln Ser Pro Glu Gly Cys Ser Ser Tyr Thr Phe Pro Lys Ile Met
 245 250 255
 Ser Pro Ala Lys Ala Thr Glu Met Leu Ile Phe Gly Lys Lys Leu Thr
 260 265 270
 Ala Gly Glu Ala Cys Ala Gln Gly Leu Val Thr Glu Val Phe Pro Asp
 275 280 285
 Ser Thr Phe Gln Lys Glu Val Trp Thr Arg Leu Lys Ala Phe Ala Lys
 290 295 300
 Leu Pro Pro Asn Ala Leu Arg Ile Ser Lys Glu Val Ile Arg Lys Arg
 305 310 315 320

Glu Arg Glu Lys Leu His Ala Val Asn Ala Glu Glu Cys Asn Val Leu
 325 330 335

Gln Gly Arg Trp Leu Ser Asp Glu Cys Thr Asn Ala Val Val Asn Phe
 340 345 350

Leu Ser Arg Lys Ser Lys Leu
 355

<210> 107
 <211> 530
 <212> PRT
 <213> Homo sapiens

<400> 107
 Met Phe Gln Phe His Ala Gly Ser Trp Glu Ser Trp Cys Cys Cys Cys
 1 5 10 15

Leu Ile Pro Ala Asp Arg Pro Trp Asp Arg Gly Gln His Trp Gln Leu
 20 25 30

Glu Met Ala Asp Thr Arg Ser Val His Glu Thr Arg Phe Glu Ala Ala
 35 40 45

Val Lys Val Ile Gln Ser Leu Pro Lys Asn Gly Ser Phe Gln Pro Thr
 50 55 60

Asn Glu Met Met Leu Lys Phe Tyr Ser Phe Tyr Lys Gln Ala Thr Glu
 65 70 75 80

Gly Pro Cys Lys Leu Ser Arg Pro Gly Phe Trp Asp Pro Ile Gly Arg
 85 90 95

Tyr Lys Trp Asp Ala Trp Ser Ser Leu Gly Asp Met Thr Lys Glu Glu
 100 105 110

Ala Met Ile Ala Tyr Val Glu Glu Met Lys Lys Ile Ile Glu Thr Met
 115 120 125

Pro Met Thr Glu Lys Val Glu Glu Leu Leu Arg Val Ile Gly Pro Phe
 130 135 140

Tyr Glu Ile Val Glu Asp Lys Lys Ser Gly Arg Ser Ser Asp Ile Thr
 145 150 155 160

Ser Val Arg Leu Glu Lys Ile Ser Lys Cys Leu Glu Asp Leu Gly Asn
 165 170 175

Val Leu Thr Ser Thr Pro Asn Ala Lys Thr Val Asn Gly Lys Ala Glu
 180 185 190

Ser Ser Asp Ser Gly Ala Glu Ser Glu Glu Glu Glu Ala Gln Glu Glu
 195 200 205

Val Lys Gly Ala Glu His Ser Asp Asn Asp Lys Lys Met Met Lys Lys
 210 215 220

Ser Ala Asp His Lys Asn Leu Glu Val Ile Val Thr Asn Gly Tyr Asp
 225 230 235 240
 Lys Asp Gly Phe Val Gln Asp Ile Gln Asn Asp Ile His Ala Ser Ser
 245 250 255
 Ser Leu Asn Gly Arg Ser Thr Glu Glu Val Lys Pro Ile Asp Glu Asn
 260 265 270
 Leu Gly Gln Thr Gly Lys Ser Ala Val Cys Ile His Gln Gly Ile Asn
 275 280 285
 Asp Asp His Val Glu Asp Val Thr Gly Ile Gln His Leu Thr Ser Asp
 290 295 300
 Ser Asp Ser Glu Val Tyr Cys Asp Ser Met Glu Gln Phe Gly Gln Glu
 305 310 315 320
 Glu Ser Leu Asp Ser Phe Thr Ser Asn Asn Gly Pro Phe Gln Tyr Tyr
 325 330 335
 Leu Gly Gly His Ser Ser Gln Pro Met Glu Asn Ser Gly Phe Arg Glu
 340 345 350
 Asp Ile Gln Val Pro Pro Gly Asn Gly Asn Ile Gly Asn Met Gln Val
 355 360 365
 Val Ala Val Glu Gly Lys Gly Glu Val Lys His Gly Gly Glu Asp Gly
 370 375 380
 Arg Asn Asn Ser Gly Ala Pro His Arg Glu Lys Arg Gly Gly Glu Thr
 385 390 395 400
 Asp Glu Phe Ser Asn Val Arg Arg Gly Arg Gly His Arg Met Gln His
 405 410 415
 Leu Ser Glu Gly Thr Lys Gly Arg Gln Val Gly Ser Gly Gly Asp Gly
 420 425 430
 Glu Arg Trp Gly Ser Asp Arg Gly Ser Arg Gly Ser Leu Asn Glu Gln
 435 440 445
 Ile Ala Leu Val Leu Met Arg Leu Gln Glu Asp Met Gln Asn Val Leu
 450 455 460
 Gln Arg Leu Gln Lys Leu Glu Thr Leu Thr Ala Ala Lys Ser Ser Thr
 465 470 475 480
 Ser Thr Leu Gln Thr Ala Pro Gln Pro Thr Ser Ser Gln Arg Pro Ser
 485 490 495
 Trp Trp Pro Phe Glu Met Ser Pro Gly Val Leu Thr Phe Ala Ile Ile
 500 505 510
 Trp Pro Phe Ile Ala Gln Trp Leu Val Tyr Leu Tyr Tyr Gln Arg Arg
 515 520 525

Arg Arg
530

<210> 108
<211> 20
<212> PRT
<213> Homo sapiens

<400> 108
Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu Asp
1 5 10 15

Phe Thr Gly Lys
20

<210> 109
<211> 20
<212> PRT
<213> Homo sapiens

<400> 109
Arg Ala Thr Val Gly Asn Ile Lys Thr Glu Arg Pro Gly Met Val Asp
1 5 10 15

Phe Lys Gly Lys
20

<210> 110
<211> 20
<212> PRT
<213> Homo sapiens

<400> 110
Gln Ala Val Ile Gly Asn Ile Asn Ile Glu Cys Ser Glu Met Leu Glu
1 5 10 15

Leu Lys Gly Lys
20

<210> 111
<211> 20
<212> PRT
<213> Homo sapiens

<400> 111
Gln Ala Ile Ile Gly Asp Ile Asn Ile Glu Tyr Leu Gly Met Leu Asp
1 5 10 15

Phe Lys Gly Lys
20

<210> 112
<211> 20
<212> PRT
<213> Homo sapiens

<400> 112
Gln Ala Ile Val Gly Asp Ile Asn Ile Ala Cys Pro Gly Met Leu Asp
1 5 10 15

Leu Lys Gly Lys
20

<210> 113
<211> 20
<212> PRT
<213> Homo sapiens

<400> 113
Gln Ala Thr Val His Asp Leu Asn Thr Glu Trp Pro Arg Met Leu Asp
1 5 10 15

Leu Lys Gly Lys
20

<210> 114
<211> 20
<212> PRT
<213> Homo sapiens

<400> 114
Gln Val Lys Val Gly Asn Cys Asn Thr Pro Lys Pro Ser Phe Phe Asp
1 5 10 15

Phe Glu Gly Lys
20

<210> 115
<211> 20
<212> PRT
<213> Homo sapiens

<400> 115
Gln Ala Thr Glu Gly Pro Cys Asn Met Pro Lys Pro Gly Val Phe Asp
1 5 10 15

Leu Ile Asn Lys
20

<210> 116
<211> 20
<212> PRT
<213> Homo sapiens

<400> 116
Gln Ala Thr Glu Gly Pro Cys Lys Leu Ser Arg Pro Gly Phe Trp Asp
1 5 10 15

Pro Ile Gly Arg
20

<210> 117
<211> 20
<212> PRT
<213> Homo sapiens

<400> 117
Gln Ala Thr Gln Gly Asp Cys Asp Ile Pro Gly Pro Pro Ala Ser Asp
1 5 10 15

Val Arg Ala Arg
20

<210> 118
<211> 18
<212> PRT
<213> Homo sapiens

<400> 118
Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu Asp
1 5 10 15

Phe Thr

<210> 119
<211> 18
<212> PRT
<213> Homo sapiens

<400> 119
Gln Ala Thr Val Gly Asp Val Asn Thr Asp Arg Pro Gly Leu Leu Asp
1 5 10 15

Leu Lys

<210> 120
<211> 18
<212> PRT
<213> Homo sapiens

<400> 120
Arg Ala Thr Val Gly Asn Ile Lys Thr Glu Arg Pro Gly Met Val Asp
1 5 10 15

Phe Lys

<210> 121
 <211> 32
 <212> PRT
 <213> Bos taurus

<400> 121
 Ile Tyr Ser His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
 1 5 10 15
 Arg Pro Gly Met Leu Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp
 20 25 30

<210> 122
 <211> 32
 <212> PRT
 <213> Homo sapiens

<400> 122
 Ile Tyr Gly His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
 1 5 10 15
 Arg Pro Gly Met Leu Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp
 20 25 30

<210> 123
 <211> 32
 <212> PRT
 <213> Drosophila melanogaster

<400> 123
 Leu Tyr Ser Leu Tyr Lys Gln Ala Thr Val Gly Asp Cys Asn Thr Asp
 1 5 10 15
 Lys Pro Gly Phe Leu Asp Phe Lys Gly Lys Ala Lys Trp Glu Ala Trp
 20 25 30

<210> 124
 <211> 32
 <212> PRT
 <213> Gallus gallus

<400> 124

Val Tyr Ser His Tyr Lys Gln Ala Thr Val Gly Asp Val Asn Thr Asp
 1 5 10 15

Arg Pro Gly Met Leu Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp
 20 25 30

<210> 125

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic
 construct; chemically synthesized

<400> 125

Ile Tyr Ser His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
 1 5 10 15

Arg Pro Gly Met Leu Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp
 20 25 30

<210> 126

<211> 32

<212> PRT

<213> Homo sapiens

<400> 126

Ile Tyr Gly His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
 1 5 10 15

Arg Pro Gly Met Leu Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp
 20 25 30

<210> 127

<211> 32

<212> PRT

<213> turtle

<400> 127

Ile Tyr Ser His Phe Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
 1 5 10 15

Arg Pro Gly Phe Leu Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp
 20 25 30

<210> 128
<211> 32
<212> PRT
<213> mallard

<400> 128
Val Tyr Ser His Tyr Lys Gln Ala Thr Val Gly Asp Val Asn Thr Asp
1 5 10 15
Arg Pro Gly Met Leu Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp
20 25 30

<210> 129
<211> 32
<212> PRT
<213> Mus musculus

<400> 129
Ile Tyr Ser His Phe Lys Gln Ala Thr Val Gly Asp Val Asn Thr Asp
1 5 10 15
Arg Pro Gly Leu Leu Asp Leu Lys Gly Lys Ala Lys Trp Asp Ser Trp
20 25 30

<210> 130
<211> 32
<212> PRT
<213> Sus scrofa

<400> 130
Ile Tyr Ser His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
1 5 10 15
Arg Pro Gly Ile Leu Asp Leu Lys Gly Lys Ala Lys Trp Asp Ala Trp
20 25 30

<210> 131
<211> 32
<212> PRT
<213> Bos taurus

<400> 131

Ile Tyr Ser His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
1 5 10 15

Arg Pro Gly Met Leu Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp
20 25 30

<210> 132

<211> 32

<212> PRT

<213> Homo sapiens

<400> 132

Ile Tyr Gly His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
1 5 10 15

Arg Pro Gly Met Leu Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp
20 25 30

<210> 133

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic
construct; chemically synthesized

<400> 133

Ile Tyr Ser His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
1 5 10 15

Arg Pro Gly Met Leu Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp
20 25 30

<210> 134

<211> 32

<212> PRT

<213> Homo sapiens

<400> 134

Ile Tyr Gly His Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
1 5 10 15

Arg Pro Gly Met Leu Asp Phe Thr Gly Lys Ala Lys Trp Asp Ala Trp
 20 25 30

<210> 135
 <211> 32
 <212> PRT
 <213> Anas platyrhynchos

<400> 135
 Leu Tyr Gly Phe Tyr Lys Gln Ala Thr Val Gly Asp Ile Asn Ile Glu
 1 5 10 15

Cys Pro Gly Met Leu Asp Leu Lys Gly Lys Ala Lys Trp Glu Ala Trp
 20 25 30

<210> 136
 <211> 32
 <212> PRT
 <213> turtle

<400> 136
 Ile Tyr Ser His Phe Lys Gln Ala Thr Val Gly Asp Ile Asn Thr Glu
 1 5 10 15

Arg Pro Gly Phe Leu Asp Phe Lys Gly Lys Ala Lys Trp Asp Ala Trp
 20 25 30

<210> 137
 <211> 20
 <212> PRT
 <213> Homo sapiens

<400> 137
 Gln Ser Thr Val Gly Asp Ile Asn Ile Glu Cys Pro Gly Met Leu Asp
 1 5 10 15

Leu Lys Gly Lys
 20

<210> 138
 <211> 20
 <212> PRT
 <213> Homo sapiens

<400> 138
 Gln Ala Ser Val Gly Asp Asn Asp Thr Ala Lys Pro Gly Leu Leu Asp
 1 5 10 15

Leu Lys Gly Lys
 20

<210> 139
 <211> 20
 <212> PRT
 <213> Homo sapiens

<400> 139
 Gln Ala Ser Val Gly Asp Asn Asp Thr Ala Lys Pro Gly Leu Leu Asp
 1 5 10 15

Leu Lys Gly Lys
 20

<210> 140
 <211> 20
 <212> PRT
 <213> Homo sapiens

<400> 140
 Gln Ala Thr Val Gly Asp Asn Asn Thr Glu Lys Pro Gly Leu Leu Asp
 1 5 10 15

Leu Lys Gly Lys
 20

<210> 141
 <211> 20
 <212> PRT
 <213> Bos taurus

<400> 141
 Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Met Leu Asp
 1 5 10 15

Phe Lys Gly Lys
 20

<210> 142
 <211> 20
 <212> PRT
 <213> Mus musculus

<400> 142
 Gln Ala Thr Val Gly Asp Val Asn Thr Asp Arg Pro Gly Leu Leu Asp
 1 5 10 15

Leu Lys Gly Lys

20

<210> 143
<211> 20
<212> PRT
<213> Rattus norvegicus

<400> 143
Gln Ala Thr Val Gly Asp Val Asn Thr Asp Arg Pro Gly Leu Leu Asp
1 5 10 15
Leu Lys Gly Lys
20

<210> 144
<211> 20
<212> PRT
<213> Sus scrofa

<400> 144
Gln Ala Thr Val Gly Asp Ile Asn Thr Glu Arg Pro Gly Ile Leu Asp
1 5 10 15
Leu Lys Gly Lys
20

<210> 145
<211> 20
<212> PRT
<213> Bos taurus

<400> 145
Gln Ala Thr Glu Gly Pro Cys Lys Leu Ser Lys Pro Gly Phe Trp Asp
1 5 10 15
Pro Val Gly Arg
20

<210> 146
<211> 20
<212> PRT
<213> Cyprinus carpio

<400> 146
Gln Ala Thr Gln Gly Pro Cys Asn Thr Pro Lys Pro Ser Met Leu Asp
1 5 10 15
Phe Val Asn Lys
20

<210> 147
<211> 20

<212> PRT
 <213> Mus musculus

 <400> 147
 Gln Ala Thr Glu Gly Thr Cys Asn Met Pro Lys Pro Gly Met Leu Asp
 1 5 10 15

 Phe Val Asn Lys
 20

 <210> 148
 <211> 20
 <212> PRT
 <213> Homo sapiens

 <220>
 <221> VARIANT
 <222> (2)
 <223> wherein Xaa is any amino acid

 <220>
 <221> VARIANT
 <222> (3)
 <223> wherein Xaa is any amino acid

 <220>
 <221> VARIANT
 <222> (6)
 <223> wherein Xaa is any amino acid

 <220>
 <221> VARIANT
 <222> (7)
 <223> wherein Xaa is any amino acid

 <220>
 <221> VARIANT
 <222> (10)
 <223> wherein Xaa is any amino acid

 <220>
 <221> VARIANT
 <222> (11)
 <223> wherein Xaa is Arg or Lys

 <220>
 <221> VARIANT
 <222> (13)
 <223> wherein Xaa is any amino acid

 <220>
 <221> VARIANT
 <222> (14)
 <223> wherein Xaa is any amino acid

 <220>

<221> VARIANT
 <222> (15)
 <223> wherein Xaa is any amino acid

<220>
 <221> VARIANT
 <222> (18)
 <223> wherein Xaa is any amino acid

<400> 148
 Gln Xaa Xaa Val Gly Xaa Xaa Asn Thr Xaa Xaa Pro Xaa Xaa Xaa Asp
 1 5 10 15
 Phe Xaa Gly Lys
 20

<210> 149
 <211> 89
 <212> PRT
 <213> Homo sapiens

<400> 149
 Thr Ala Ser Thr Thr Pro Cys Ala Lys Trp Ser Ser Ser Cys Ala Ala
 1 5 10 15
 Leu Lys Gln Leu Lys Gly Pro Val Ser Asp Gln Glu Lys Leu Leu Val
 20 25 30
 Tyr Gly Leu Tyr Lys Gln Ala Thr Gln Gly Asp Cys Asp Ile Pro Gly
 35 40 45
 Pro Pro Ala Ser Asp Val Arg Ala Arg Ala Lys Trp Glu Ala Trp Ser
 50 55 60
 Ala Asn Lys Gly Ala Ser Lys Met Asp Ala Met Arg Gly Tyr Ala Ala
 65 70 75 80
 Lys Val Glu Glu Leu Thr Lys Lys Glu
 85

<210> 150
 <211> 228
 <212> PRT
 <213> Homo sapiens

<400> 150
 Met Gly Asp Ala Gly Ala Thr Ala Ala Ala Leu Arg Pro Ala His Asn
 1 5 10 15
 Leu Arg Pro Ala Pro Pro Thr Ala Ser Ala Ala His Ala Gln Ser Ser
 20 25 30
 Arg Thr Ser Ala Pro Ser Ala Gln Arg Arg Leu Pro Ala Glu Pro Ser
 35 40 45

His Gln Pro Ser Ala Pro Gly Thr Ala Ser Thr Thr Pro Cys Ala Lys
 50 55 60
 Trp Ser Ser Ser Cys Ala Ala Leu Lys Gln Leu Lys Gly Pro Val Ser
 65 70 75 80
 Asp Gln Glu Lys Leu Leu Val Tyr Gly Leu Tyr Lys Gln Ala Thr Gln
 85 90 95
 Gly Asp Cys Asp Ile Pro Gly Pro Pro Ala Ser Asp Val Arg Ala Arg
 100 105 110
 Ala Lys Trp Glu Ala Trp Ser Ala Asn Lys Gly Ala Ser Lys Met Asp
 115 120 125
 Ala Met Arg Gly Tyr Ala Ala Lys Val Glu Glu Leu Thr Lys Lys Glu
 130 135 140
 Val Gly Gly Val Glu Arg Glu Gln Arg Gly Val Gln Asp Gly Arg His
 145 150 155 160
 Glu Gly Leu Arg Gly Gln Ser Gly Gly Ala Asp Glu Glu Gly Arg Ala
 165 170 175
 Ser Lys Met Asp Ala Met Arg Gly Tyr Ala Ala Lys Val Glu Glu Leu
 180 185 190
 Thr Lys Lys Glu Val Gly Gly Val Glu Arg Glu Gln Arg Gly Val Gln
 195 200 205
 Asp Gly Arg His Glu Gly Leu Arg Gly Gln Ser Glu Glu Met Arg Lys
 210 215 220
 Lys Glu Ala Gly
 225

<210> 151
 <211> 191
 <212> PRT
 <213> Homo sapiens

<400> 151
 Met Gly Asp Ala Gly Ala Thr Ala Ala Ala Leu Arg Pro Ala His Asn
 1 5 10 15
 Leu Arg Pro Ala Pro Pro Thr Ala Ser Ala Ala His Ala Ser Pro His
 20 25 30
 Glu Arg Ala Arg Gln Ala Ser Arg Ala Phe Arg Gln Ser Pro Pro Thr
 35 40 45
 Ser Pro Gln Leu Leu Ala Pro Gly Thr Ala Ser Thr Thr Pro Cys Ala
 50 55 60
 Lys Trp Ser Ser Ser Cys Ala Ala Leu Lys Gln Leu Lys Gly Pro Val
 65 70 75 80

Ser Asp Gln Glu Lys Leu Leu Val Tyr Gly Leu Tyr Lys Gln Ala Thr
 85 90 95

Gln Gly Asp Cys Asp Ile Pro Gly Pro Pro Ala Ser Asp Val Arg Ala
 100 105 110

Arg Ala Lys Trp Glu Ala Trp Ser Ala Lys Lys Gly Ala Ser Lys Met
 115 120 125

Asp Ala Met Arg Gly Tyr Ala Ala Lys Val Glu Glu Leu Thr Lys Lys
 130 135 140

Glu Val Gly Gly Val Glu Arg Glu Gln Arg Gly Val Gln Asp Gly Arg
 145 150 155 160

His Glu Gly Leu Arg Gly Gln Ser Gly Gly Ala Asp Glu Glu Gly Ser
 165 170 175

Gly Gly Arg Gly Ala Arg Thr Lys Gly Arg Pro Arg Trp Thr Pro
 180 185 190